#### STIC-Biot ch/ChemLib

From: Sent:

Schultz, James

Wednesday, May 07, 2003 9:22 AM STIC-Biotech/ChemLib

To: Subject:

Sequence search 10/057,550

MAY -7 2003

LECH/CHEM. DIVISION (STIC)

Whoops, sorry, the corrected file name is now listed in the title above. .

----Original Message-----

STIC-Biotech/ChemLib

From: Sent:

Wednesday, May 07, 2003 9:06 AM

To:

Schultz, James

Subject:

RE: Sequence search pct/us02/04123

This Application Number does not have a valid CRF - please provide the necessary correction or another application

maude

-----Original Message-----Schultz, James

From: Sent:

Wednesday, May 07, 2003 8:36 AM

To:

STIC-Biotech/ChemLib

Subject: Sequence search pct/us02/04123

Hello,

I need a length limited nucleotide sequence search performed on SEQ ID NO: 64 in the above entitled application, where the maximum size of the returned hit is no more than 50 nucleotides long. Thanks, Doug Schultz

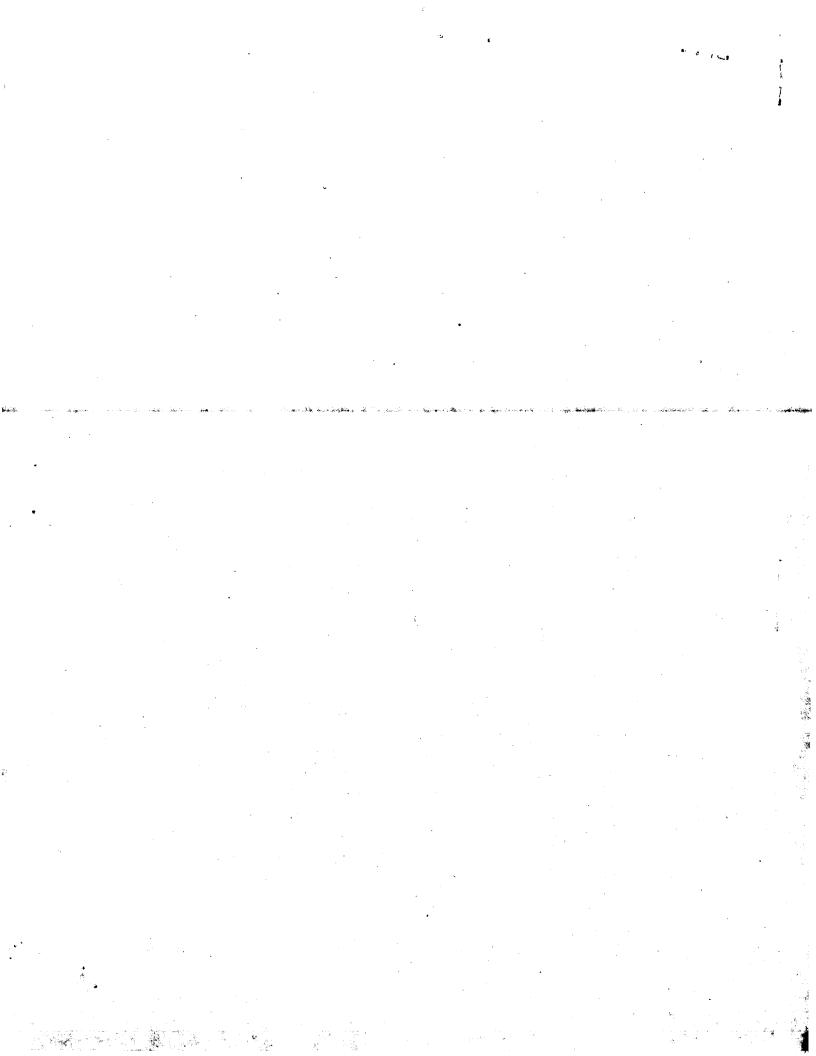
J. Douglas Schultz, Ph.D. AU 1635 (Biotechnology) Patent Examiner United States Patent and Trademark Office (703) 308-9355 (703) 746-3973 (fax) Office: CM1 12E18 Mail: CM1 11E12

Edward Hart Technicai Info. Specialist STIC/Biotech CMI 6B02 Tel: 305-9203

Searcher:
Phone:
Location:
Date Picked Up: 518/105
Date Completed: 5/17/1/35
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH;
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (where app	olic.)
STN:	
DIALOG:	
Questel/Orbit:	
DRLink:	
Lexis/Nexis:	
Sequence Sys.: / tex 10/1	
WWW/Internet:	
Other (specify):	





#### STIC SEARCH RESULTS

#### Biotech-Chem Library

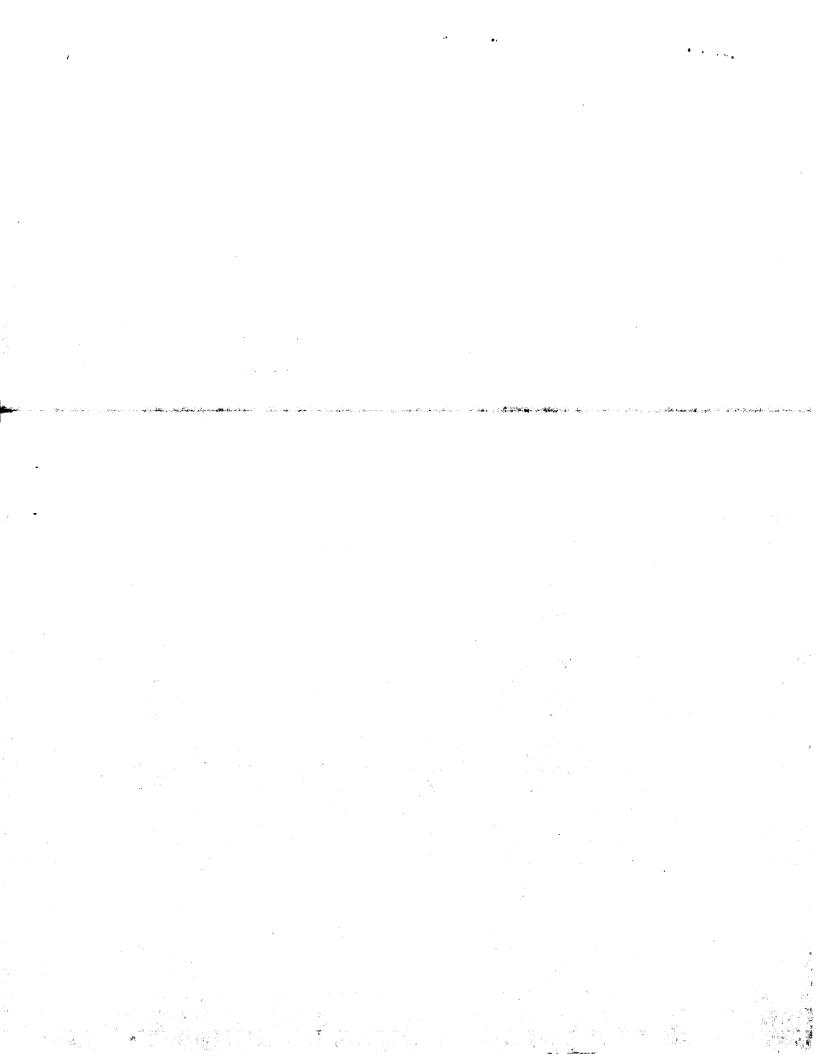
Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 308-4258, CM1-1E01

70l	untary Results Feedback Full
>	I am an examiner in Workgroup: Example: 1610
>	Relevant prior art found, search results used as follows:
	☐ 102 rejection
	☐ 103 rejection
	Cited as being of interest.
	Helped examiner better understand the invention.
	Helped examiner better understand the state of the art in their technology.
	Types of relevant prior art found:
	☐ Foreign Patent(s)
	Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
>	Relevant prior art not found:
	Results verified the lack of relevant prior art (helped determine patentability).
	Results were not useful in determining patentability or understanding the invention.
Со	mments:

Diopoilor sand completed forms to STECE beak.





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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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## SUMMARIES

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### ALIGNMENTS

REFERENCE AUTHORS		ORGANISM	SOURCE	SEGMENT	KEYWORDS	VERSION	ACCESSION ·	DEFINITION	LOCUS	RESULT 1 HUMRAF12
<pre>1 (bases: 1 to 49) Bonner,T.I., Kerby,S.B., Sutrave,P., Gunnell,M.A., Mark,G. and Rapp,U.R.</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Homo sapiens	Homo sapiens (tissue library: of Lawn et al.) DNA.	2 of 9	<pre>c-myc proto-oncogene; raf protein; raf proto-oncogene.</pre>	L00213.1 GI:190830	L00213 ·	Human c-raf-1 proto-oncogene, exon 2, clones lambda-(2:13).	HUMRAF12 49 bp DNA linear PRI 08-JAN-1995	

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Patent: WO 0166800-A 835 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Mammalla; Eutheria; Primates; Catarrhini 1 (bases 1 to 31)
Cargill,M., Ireland,J.S. and Lander,E.S.
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Mol. Cell. Biol. 5 (6), 1400-1407 (1985)
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1 (bases 1 to 3
Chenchik, A., Jo)
                                                    1 (bases 1 to 26)
Chenchik, A. Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 5994076-A 980 30-NOV-1999;
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Patent: US 6352829-A 979 05-MAR-2002;
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Chenchik, A., Jokha
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cooperating oncogenes

Length 26;

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196169
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1 (bases 1 to 26)

1 (bases 1 to 26)

Chenchik, A., Jokhadze, G. and Bibilashvilli, R. Methods of assaying differential expression Patent: US 6352829-A 980 05-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
AR197895
                                                                                                                                                                                                   Unclassified.

1 (bases 1 to 26)

Calabretta,B. and Skorski,T.

Antisense oligonucleotides targeting cooperating oncogenes

Patent: US 5734039-A 5 31-MAR-1998;
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l Similarity 100.0%;
26; Conservative
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196168
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26; Conser
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4 c 14 g
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6 c 9 g
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AR110776
                          Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucle
methods of use thereof
Patent: WO 0140521-A 485 07-JUN-2001;
Curagen Corporation (US)
                                                                                                                                                   Sequence 485
AX157157
                                                                                                                                                                                                                                                                                                                                                        Kasid, U., Gokhale, P., Dritschilo, A. a Liposomes containing oligonucleotides Patent: US 6126965-A 2 03-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Calabretta,B. and Skorski,T. Antisense oligonucleotides targetin Patent: US 5734039-A 6 31-MAR-1998;
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Unclassified
                                                                                                           Homo sapiens
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7 c 6 g
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/organism="Homo sapiens"
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22-JUN-2001

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                       28;
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Pavlakis,G.N. and Felber,B.K.
Nucleic acid constructs containing H
inhibitory/instability regions and m
Patent: US 5972596-A 17 26-OCT-1999;
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Sequence
AR079693
                                                                                                                                                                                                                                                                                                                              ch 0.7%;
l Similarity 73.7%;
28; Conservative
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AR081223
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Method of eliminating inhibitory/ instability
Patent: US 5965726-A 17 12-OCT-1999;
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                                Similarity
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Accession number cg44928329"
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                    Score 22; DB Pred. No. 1.5e 0; Mismatches
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Pred. No. 1.3e+
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RESULT 15
AX164948/c
LOCUS
                                        VERSION
KEYWORDS
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FEATURES
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SOURCE
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AR170583
LOCUS
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                       ORGANISM
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Sequence 14
AX164948
AX164948.1
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Sequence
AR170583
 Eukaryota; Metazoa;
Mammalia; Eutheria;
                      Homo sapiens
                                human.
                                                                                                                                                                                                                                                          1 (bases 1 to 43)
Pavlakis,G.N. and Felber,B.K.
Patchod of eliminating inhibitory/instability
Patent: US 6291664-A 17 18-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AR142002
Sequence
AR142002
                                                                                                                                                                          28; Conserv
                                                                                                                                                                                                                                                                                                                  Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 73.7
28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pavlakis,G.N. and Felber,B.K. Method of eliminating inhibitory/instability Patent: US 6174666-A 17 16-JAN-2001;
                                                                                                                                                                                                                                                                                                        Unclassified.
                                                                                                                                                                                                                                                                                                                            Unknown.
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11 c 5 q
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                                                   GI:14545777
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from patent US 6291664.
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Chordata;
Primates;
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                                                                                                                                                                        Score 22; DB
Pred. No. 1.5e
0; Mismatches
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Pred. No. 1.5e+
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Craniata; Vo
Catarrhini;
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1.5e+06;
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          Vertebrata;
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         Euteleostomi;
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Hominidae;

Homo

GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993. - 2003 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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2977
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                      Length
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*
                                                                                                                                                                      DB
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AAA07016
AAA07027
AAA3030347
AAI30349
AAK66891
AAV20480
ABK66892
AAL31396
AAH444455
                                                                                                                                                                                                                                                                                                                                       geneseq/qeneseqn-embl
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                                                                                                                                                                    Description
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Primer KASSIIA		AAVUS431	19	4		21	<b>4</b> U
		AAV05431	19	40	0.7	21	5
Human c-Raf		ABL50208	24	27	0.7	21	44
Human c-Raf		ABL50176	24	27	0.7	21	43
Human c-Raf		AAH41589	22	27	0.7	21	42
Human gene		AAF97098	22	21	0.7	21	41
Antitumoural	٨	AAT93823	18	35	0.7	$\vdash$	40
GAD plasmid		AAA55866	21	40	0.7	21.4	39
RAF PCR primer		AAA55851	21	40	0.7	21.4	38
Yeast PCR		AAA72893	21	40	0.7	21.4	37
Yeast RAF		AAA72878	21	40	0.7	21.4	36
Primer used		AAV05418	19	40	0.7	21.4	35
Primer RASSfiS		AAV05430	1.9	40	0.7	_	34
Human sile		AAI75927	22	50	0.7	1.	
Human silent		ABL00152	23	50	0.7	22	
		AAZ69520	21	47	0.7	22	
c-fas position		AAQ50213	14	43	0.7	22	
Phosphodiester 4 1 2		AAT93841	18	41	0.7	22	29
. Human c-Raf1		ABL50209	24	30	0.7	22	
Human c-Rafl		ABL50177	24	30	0.7	22	27
		AAH41590	22	30	0.7	22	
Human silent		AAI73544	22	50	0.7	2	
Phosphodiester ol		AAT93834	18	27	0.7	22.2	
Human SNP		AAL31673	22	50	•	2	
Human c-Raf		ABL50229	24	27	•	2	22
Human c-Rafl		ABL50197	24	27		22.4	21
an		AAH41610	22	27	0.8	ν.	20
C-raf antisense		AAA26246	21	23	•	23	19
C-raf antisens		AAA26245	21	23	٠	23	
C-raf antisens		AAA26244	21	23.	٠	23	17
-raf ant		AAA26243	21	23	٠	23	16
C-raf antisense		AAA26242	21	23	٠	23	15
C-raf antisense	•	AAA26239	21	23	0.8	23	14
Hairpin hy		AAA29998	21	ω G	0.8	24	13
5		AAA30000	21	31	0.8	24	12
C-raf probe		30	19	24	0.8	24	11
Hairpin hybridizer		AAA29996	21	35	0.8	24.2	10

### ALIGNMENTS

03-JUL-2000 AAA07016; AAA07016 standard; DNA; 42 BP

RESULT 1 AAA07016

Raf-1 mutagenic PCR primer, SEQ ID NO:13.

(first entry)

sphingomyelin mutagenesis; lipopolysaccharide; Raf-1; CAP kinase; phosphorylation; ceramide-activated protein kinase; endotoxin; PCR primer;

Synthetic. Mammalia.

US6040149-A.

21-MAR-2000

10-JAN-1997; 97US-0785247.

11-JAN-1996; 96US-0009900.

(SLOK ) SLOAN KETTERING INST CANCER RES

Liu J, Kolesnick RN;

WPI; 2000-270133/23.

latexin

Novel method of identifying agents capable of inhibiting

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RESULT 2
AAA07027
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Best Local S
Matches 38
WPI; 2000-270133/23
                                                    Zhang Y, Liu J,
                                                                                                 (SLOK ) SLOAN KETTERING INST CANCER RES
                                                                                                                                                           11-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    weight of 100-110 kD. It is an upstream participant in a sphingomyelin signal transduction pathway which uses ceramide as a second messenger. This pathway is initiated by tumour necrosis factor-alpha (TNF-alpha) and interleukin-beta (II-beta), causing the hydrolysis of sphingomyelin to ceramide. The ceramide in turn stimulates the kinase to phosphorylate protein substrates which can then mediate signal transduction. The CAP kinase is also stimulated by the bacterial endotoxin lipopolysaccharide (IPS), which is thought to mimic the second messenger function of ceramide. The methods are useful for identifying agents that inhibit lipopolysaccharide-induced Thr phosphorylation by CAP kinase. The agents identified using themethod are useful for
                                                                                                                                                                                                             10-JAN-1997;
                                                                                                                                                                                                                                                                     21-MAR-2000
                                                                                                                                                                                                                                                                                                                          US6040149-A
                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Raf-1; CAP kinase; phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Raf-1 mutagenic PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA07027;
                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             sphingomyelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mutagenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA07027 standard; DNA; 39 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    erythematosus, HIV, infection, disorders associated with poor stream of the septic shock. Sequences AAA07016-A07019 and AAA07027 represent PCR primers used in an exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treating disorders associated with aberrant phosphorylation of target molecules by CAP kinase, e.g., inflammatory disorders (such as rheumatoid arthritis), ulcerative colitis, graft versus host disease, lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method of determining whether an agent is capable of specifically inhibiting the ability of a ceramide-activated protein (CAP) kinase to phosphorylate the threonine residue in a polypeptide containing a Thr-Pro- or Thr-Leu-Pro motif. In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thereof.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 9 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                  signal transduction pathway; PCR primer;
                                                                                                                                                           96us-0009900
                                                                                                                                                                                                             97US-0785247
                                                  Kolesnick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Flag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.2%;
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ceramide-activated protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Sequence

39 BP;

9

Α,

11 C; 13

G; 6 T; 0 other;

0,

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phosphorylate protein substrates which can then mediate signal transduction. The CAP kinase is also stimulated by the bacterial endotoxin lipopolysaccharide (LPS), which is thought to mimic the second messenger function of ceramide. The methods are useful for identifying agents that inhibit lipopolysaccharide-induced Thr phosphorylation by CAP kinase. The agents identified using the method are useful for molecules by CAP kinase, e.g., inflammatory disorders (such as rheumatoid arthritis), ulcerative colitis, graft versus host disease, lupus erythematosus, HTV, infection, disorders associated with poor stem cell growth, and septic shock. Sequences AAA07016-A07019 and AAA07027 represent PCR primers used in an exemplification of the present invention to generate mutant Flag peptide/Raf-1 sequences via overlap extension
                                                                                                                                                                                                                                                                                                                                                                                                 weight of 100-110 kD. It is an upstream participant in a sphingomyelin signal transduction pathway which uses ceramide as a second messenger. This pathway is initiated by tumour necrosis factor-alpha (TNF-alpha) and interleukin beta (IL-beta), causing the hydrolysis of sphingomyelin to ceramide. The ceramide in turn stimulates the kinase to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel method of deterministic capable of specifically inhibiting the ability of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel method of identifying agents capable of inhibiting lipopolysaccharide induced threonine phosphorylation by ceramide-activated protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         residue in a polypeptide containing a Thr-Pro- or Thr-Leu-Pro motif. In particular, the peptide substrate that is specifically phosphorylated is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ceramide-activated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Raf-1, epidermal growth factor receptor (EGFR), or suitable fragments thereof. The CAP kinase is membrane bound and has an apparent molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example VI; Column 56; 84pp; English
                                                         Flag peptide/Raf-1 sequences via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CAP) kinase to phosphorylate the threonine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of determining whether an agent
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RESULT 3 AAA07017 BX 8X H В δÃ Matches Query Match Zhang Y, (SLOK ) SLOAN KETTERING INST CANCER RES 11-JAN-1996; 10-JAN-1997; 21-MAR-2000 US6040149-A Synthetic. Mammalia. sphingomyelin mutagenesis; Raf-1; CAP kinase; phosphorylation; ceramide-activated protein kinase; Raf-1 mutagenic PCR primer, SEQ ID NO:14. 03-JUL-2000 AAA07017; AAA07017 standard; DNA; 42 919 1 CACATGGTCAGCACCGTGCTACCGGTGGACAGCAGGATG CACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG l Similarity 35; Conserv Liu (first entry) Conservative lipopolysaccharide; LPS; endotoxin; signal transduction pathway; PCR primer; ss 'n 96US-0009900 97US-0785247 Kolesnick 1.1%;
89.7%; 0 Score 32.6; Pred. No. 1: Mismatches DВ 957 39 21; 4 Length Indels 39; 0 Gaps

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RESULT 4
AA130347
ID AA13
AC AA13
AC AA13
XX Humm
CON Humm
CON Home
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method of determining whether an agent CC is capable of specifically inhibiting the ability of a CC caramide-activated protein (CRP) kinase to phosphorylate the threonine CC particular, the peptide containing a Thr-Pro- or Thr-Leu-Pro motif. In CC particular, the peptide substrate that is specifically phosphorylated is Raf-1, epidermal growth factor receptor (EGFR), or suitable fragments CC thereof. The CAP kinase is membrane bound and has an apparent molecular CC weight of 100-110 kD. It is an upstream participant in a sphingomyelin CC signal transduction pathway which uses ceramide as a second messenger. CC This pathway is initiated by tumour necrosis factor-alpha (TRP-alpha) CC and interleukin-beta (IL-beta), causing the hydrolysis of sphingomyelin CC to ceramide. The ceramide in turn stimulates the kinase to phosphorylate protein substrates which can then mediate signal CC transduction. The CAP kinase is also stimulated by the bacterial conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second conditions of coramide (LPS), which is thought to mimic the second coramic coramic conditions of coramic 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel method of identifying agents capable of inhibiting lipopolysaccharide induced threonine phosphorylation by ceramide-activated protein kinase
                                          07-MAR-2001; 2001WO-US07268
                                                                                                                          13-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI30347;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAI30347 standard; DNA; 31 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 42 BP; 8 A;
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                                                                                                                                                                                                          WO200166800-A2
                                                                                                                                                                                                                                                                                                                                                                      Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human single nucleotide polymorphism (SNP) RAF1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-270133/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGCAGGATG 957
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTCCACATGGTCAGCGTCGTGCTACCGGTGGACAGCAGGATG 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Similarity 85.7
36; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  resequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                Location/Qualifiers replace(16,T)
/*tag= a
                                                                                                                                                                                                                                                                             /standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genotype; disease; forensic; paternity testing;
polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.18;
85.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11 C; 15 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 32.4;
Pred. No. 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.
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2144 CCCTTTCTCCAGAGGCAGAACACATGTTTTC 2174

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Matches
             Query Match
Best Local
                                                                                                            The invention relates to the identification of nucleic acid molecules (AAI29513-AAI31314) from the human genome which include polymorphic sites which can predispose individuals to disease. Various genes from a number of individuals were resequenced and single nucleotide polymorphisms (SNPS) in these genes discovered. The method is useful for predicting the presence, absence or severity of a particular phenotype or disorder (e.g., diabetes) associated with a particular genotype. The nucleic acids
                                                                                                                                                                                                                                                         Nucleic acid molecules from the human genome which include polymorphic sites, useful in methods for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype .
                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                               Claim 1; Page 80; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                               Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-2000;
22-MAY-2000;
                                                       Sequence 31
                                                                                    testing.
                                                                                                containing the
                                                                                                                                                                                                                                                                                                                                                                                         (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                   2001-522952/57
               Similarity
                                                       BP;
                                                                                                                                                                                                                                                                                                                                                              Ireland
Conservative
                                                                                                 polymorphic sites may be
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                                                                                                                                                                                                                                                                                                                                                               JS,
             1.0%;
                                                      10 C;
                                                                                                                                                                                                                                                                                                                                                               Lander ES
                                                      5 G;
0;
             Score 31;
Pred. No.
                                                      9 T;
 Mismatches
                                                       0 other;
             DB 22;
29;
                                                                                                   useful in
0,
                         Length 31:
                                                                                                   forensics
Indels
 ö
Gaps
                                                                                                 paternity
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ABK66891 RESULT Primer; ss; DNA microarray; differential expression analysis; human 02-JUL-2002 Human gene specific PCR primer ABK66891; ABK66891 standard; DNA; 28 ب CCCTTTCTCCAGAGGCAGAACACATGTTTTC (first entry) ВР #979 31

Homo sapiens

05-MAR-2002 US6352829-B1

05-JAN-1999; 99US-0225928

21-MAY-1997; 97US-085999E

(CLON-) CLONTECH LAB INC

Chenchik A, Jokhadze G, Bibilashvilli R;

WPI; 2002-314699/35.

Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers

Example 3; SEQ ID No 979; 11pp; English

The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiclogical

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RESULT 6
AAV20480/c
ID AAV204
XX AAV204
XX AAV204
XX Human
DT 17-JUN
XX Human;
KW Human;
KW Cancer
XX Synthe
OS Synthe
OS Homo s;
XX Homo s;
XX III-le
FT allele
FT Allele
FT Allele
FT ST SEP
XX IS-SEP
XX IS-SEP
XX US5734;
XX IS-SEP
XX ANTICA;
PT Calabr;
XX Antica;
PT AAV204
PT AATICA;
PT AAV204

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Best Local
                       Anticancer composition comprising two anti-sense oligo:nucleotide(s)
- targetting cytoplasmic and nuclear oncogene(s);
                                                                                                          WPI; 1998-229882/20
                                                                                                                                                                                                                                                                               15-SEP-1994;
                                                                                                                                                                                                                                                                                                                                      15-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    US5734039-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1704 CAGTITCCAGTCGGATGTCTACTCCTAT 1731
                                                                                                                                                                                                                        (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  allele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human c-raf antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV20480;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV20480 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   is preferably performed by hybridising the labeled NAS for each of the associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, and comparing the patterns for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprises producing subpopulation of labeled was for the different physiological sources, comprising the populations for each physiological source to identify differences in the population, where the comparison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          between several different physiological sources, where the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer. The method is useful for producing a sup-population of lawers NAs which is useful for analysing the differences in the RNA profiles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conditions to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ۳
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                    В,
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Lity 100.0%;
Conservative
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                                                                                                                                                                 Skorski
                                                                                                                                                                                                                                                                            94US-0306691
                                                                                                                                                                                                                                                                                                                                94US-0306691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proto-oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proto-oncogene; neoplastic disease; anticancer;
oligonucleotide; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "phosphorothioate linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ly generate sub-population of NAs, where has a sequence complementary to a distinct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Pred. No. 2.2
0; Mismatches
                    oncogene(s)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local
between several different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the populations for each physiological sources, comprising the population, where the comparison source to identify differences in the population, where the comparison is preferably performed by hybridising the labeled NAs for each of the distinct physiological sources to an array of probe NAs stably
                                                                                                                         each gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled
                                                                                                                                                                                      The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where
                                                                                                                                                                                                                                                                                                                  Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers
                                                                                                                    NAs which is useful for analysing the differences in the RNA profiles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specific for a cytoplasmic oncogene or proto-oncogene selected from ras, raf, EGP-1, c-fms, c-ros, c-ki, c-met, c-trk, c-src, c-abl, bcr-abl, c-fgr and c-yes. The second oligonucleotide is specific for a nuclear oncogene or proto-oncogene selected from myc, jun, c-ets, c-fos, c-myb, B-myb, c-rel, c-vav, c-ski, c-spi, cyclin Dl, PML/RAR alpha, AMLI/MTGB, EZA/pri and ALL-1/AF-4. The composition is used for treating cancer. The combination of antisense oligonucleotides has synergistically enhanced ability to inhibit growth of cancer cells.
                                                                                                                                                                                                                                                                                       Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-314699/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human gene specific PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK66892 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CLON-) CLONTECH LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26 BP; 5 A; 4 C; 14 G; 3 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention. The present invention describes a composition which comprises two antisense oligonucleotides. The first oligonucleotide is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents an antisense oligonucleotide from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Column
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 ATGTGACCGCCTCCCGCTCCCTCACC 30
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26; Conser
                                                                                                                                                                                                                                                                                   SEQ ID No 980; 11pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA
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RESULT 8
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Best Local :
proteins, cytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encode by them may be used in the prevention, diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                             Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1963 TCTCTACCGAAGATCAACCGGAGCGC 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention.

Note: The sequence data for this patent did not form part of the primer and provided the patent of the patent 
                                                                                                                                                      The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopolet. apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the printed specification, but was obtained format directly from USPTO at
                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2710; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can emyloid protein; angiopolettin; apoptosis related protein; cadherin; amyloid protein; angiopolettin; apoptosis; clated protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f
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Pred. No.
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mune disease; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cyclins, polymerases, treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          electronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation;
                                                                                                                                                                                                                                           angiopoietin,
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    of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diseases associated with inappropriate expression of the proteins above. Disorders that may be prevented, diagnosed and/or treated i multifactorial diseases with a genetic component, such as autoimmu diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pat
                                    infection, immunological diseases and various inflammations sequence represents a PCR primer for human latexin protein 4
                                                       The present invention describes human latexin protein 46 (I) (I) has cytostatic, antiviral, anti-HIV, immunomodulatory and antiinflammatory activities. (I) and the polynucleotide encoding it are applicable in the diagnosis and treatment of malignant tumour, haemopathy, HIV
                                                                                                                                             Human latexin protein 46 and encoded podiagnosis and treatment of malignant tunimunological diseases and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                             Human; latexin protein 46; cytostatic; antiviral; anti-HIV; immunomodulatory; antiinflammatory; diagnosis; malignant tu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 18; 39pp; Chinese.
                                                                                                                                                                                                                                                                                 29-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                         W0200149737-A1
                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
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                          used in
                                                                                                                                                                                                                                              (SHAN-)
                                                                                                                                                                                                                                                         (UYFU-)
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                                                                                                                                                                                                                                                                                                                                                                                                                    haemopathy;
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                                                                                                                                                                                              2001-432872/46
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SHANGHAI BI
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                       example from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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                       present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 26;
                                                                                                                                                                                                                                              TECHNOLOGY LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis; malignant tumour;
gical disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ņ
                                                                                                                                                        polynucleotide, applicable in tumor, hemopathy, HIV infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NO: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the proteins listed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                  ammations. The present protein 46, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diabetes,
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RESULT 10
AAA29996/C
ID AAA2999
XX AAA299
AC AAA299
XX DAAA299
XX DAAA299
XX DNA-RN
KW C'RAf;
KW G'RAf;
KW G'RAf;
KW G'RAf;
FT MISC_F
FT
                                                                                     and modulated cellular and viral processes such as splicing, editing translation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets, it analysis of mutations in diseased cell and to detect specific RNA.
                                                                                                                                                                           molecule is used in an example of the invention and targets the nucleic form of the HPH molecule binds to and blocks the function of a target nucleic form.
  greater
                           sequence
                                                              hairpin structure improves resistance to nuclease degradation
                                                                                                                                                                                                                                  A method for modulating the function of a target sequence in a comprises treatment with a hairpin hybridizer (HPH) nucleic acid such as that represented by the present sequence. The present HP molecule is used in an example of the invention and targets the
                                                                                                                                                                                                                                                                                                                                                                                                                                         Modulating target sequence in a confor identifying gene function, by
                      localization within the cell, and uptake by cell sequence that activates RNase H (for cleaving RN
                                                                                                                                                                                                                                                                                                                                                                Claim 46; Page 76; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     hybridizer nucleic acid molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hartmann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1998;
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c-raf; inhibitor; detect; nuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hairpin hybridizer molecule #1 targeting c-raf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2000
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32..35
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23..35
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14..23
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1..13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   detect; nuclease resistance; gene function;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "2'.O-methyl-ribonucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Phosphorothioate internucleotide linkage"
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                                                                                                                                                                                                                                                                                                                                                                                                                                    cell, useful e.g. therapeutically by treatment with novel hairpin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'n
sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hairpin hybridizer molecule;
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                    cleaving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 internucleotide linkage
                                                                                                                                                                                                                                                                     nucleic acid mo
                 HPH may include a its specificity is
                                                                                                            t nucleic acid, ing, editing, any, in target targets, for
                                                                                                                                                                                                                                                                   HPH
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RESULT 11
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Best Local
                                                                                                                                                                                                                        Determining relative amounts of different nucleic acids by multiple: competitive polymerase chain reaction - involves tagging target and control sequences then amplification with generic primer pair corresponding to tagging sequences, used e.g. to determine response
                                                                                                                                                                                                                                                                                                                                                                                                       RT-PCR; tagginy ....
probe; housekeeping gene;
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                                                                                                                                                                                                                                                                   WPI; 1998-447252/38.
                                                                                                                                                                                                                                                                                    Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                       Multiplex competitive PCR RT-PCR; tagging reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                C-raf probe
                                                                                                                                                                                                 Example 1;
                                                                                                                                                                                                                                                                                                    (RIBO-)
                                                                                                                                                                                                                                                                                                                     07-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                      27-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 89.7 nes 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATCAATGCTTGCACGCTGACCACGCTCGC
                                                                                                                                                                                                                                                                                                    RIBOZYME
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                                                                                                                                                                                                                                                                                                     PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9 C; 13
                                                                                                                                                                                                                                                                                                                                                                                                              reaction; MC-PCR; reverse-
competitive amplification
C-raf; hybridisation; ss.
                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
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Pred. No. 3.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 1 T; 6 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .7e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 other
                                                                                                                                                                                                                                                                                                                                                                                                                                reverse-transcriptase PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                         reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              35:
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relative to primers #1 and #2 respectively, are used as tagging primers in the tagging reaction. A forward tagging primer has a defined sequence at its 5' end (+TAG sequence) while a reverse tagging primer has a different defined sequence at its 3' end (-TAG sequence). The purpose of the tagging reaction is to introduce the two defined sequences at the correct ends of the sequence to be amplified. The competitive amplification reaction involves using a single pair of generic primers, whose sequences are complementary to the +TAG and -TAG sequences, to amplify the different products generated from the cDNAs during the tagging step. This amplification reaction is competitive due to the use of a single primer pair to the amplify the different target RNAs. Probe #5, complementary to the the region of the resulting cDNA to be amplified during the MC reaction into a double-stranded molecule. Primers #3 and #4, reaction involves a reverse-transcriptase (RT-PCR) reaction followed by a tagging reaction and a competitive amplification reaction. The RT-PCR reaction uses a primer #2 to convert target mRNA into cDNA.

Primer #1 in combination with primer #2 is then used to convert. The present invention provides a method for determining by using the multiplex competitive PCR reaction (MC-PCR). relative amounts of two target RNA being amplified, or more different nucleic acid molecules ţ This amplification the MC-PCR A MC-PCR nested to a

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RESULT 12
AAA30000/c
ID AAA300
XX
AC AAA300
XX
DT 09-AUG
XX
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the amplified product. The MC-PCR reaction can amplify one or more target mRNAs in a sample using the primer set #1-#5 for each target mRNA. In the example given, primers #1, #2, #3, #4 and probe #5 are the C-raf primers 1 (AAV33092), 2 (AAV33093), 3 (AAV33094), 4 (AAV33095) and probe 5 respectively. These primers/probes were used to illustrate the method of the invention. The method claims to allow detection of low-abundance mRNA in small samples (e.g. 10 ng is sufficient) with high precision, and uses housekeeping genes as controls for RNA input and integrity. Also, a large number of samples may be processed simultaneously, making the process suitable for high
                                                                                                                                                                                                                                                                                                                                  misc_RNA
                                                                                                                                                                                                                                                                                                                                                                            misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                          misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene expression modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA-RNA hybrid; phosphorothioate; hairpin hybridizer molecule; HPH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 7 A; 9 C; 3 G; 5 T; 0 other;
Modulating target sequence in a cell, useful e.g. therapeutically or
                                WPI; 2000-292841/25
                                                               Hartmann T,
                                                                                                                           21-SEP-1998;
                                                                                                                                                                                        30-MAR-2000.
                                                                                                                                                                                                                      WO200017346-A2
                                                                                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hairpin hybridizer molecule #5 targeting c-raf RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           throughput screening, and does not require continuous monitoring.
                                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                         20-SEP-1999;
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                                                             Zwick M,
                                                                                                                           98US-0101174
                                                                                                                                                         99WO-US21865
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21..31
                                                                                                                                                                                                                                                                                                                                                                              /note=
12..20
                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                       /note-
                                                                                                                                                                                                                                                                                                 note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
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                                                                                                                                                                                                                                                                                                                                                               *tag=
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                                                               Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                       "2'-0-methyl-ribonucleotides"
                                                                                                                                                                                                                                                                                                "2'-0-methyl-ribonucleotides
                                                                                                                                                                                                                                                                                                                                             "Phosphorothioate internucleotide linkage'
                                                                                                                                                                                                                                                                                                                                                                                           "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                    "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nuclease resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
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Pred. No.
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                                                               Jarvis
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3.3e+0
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                                                               H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene function;
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for identifying gene function, by treatment with novel hairpin hybridizer nucleic acid molecules
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analysis of mutations in diseased cell and to detect specific RNA. The hairpin structure improves resistance to nuclease degradation, localization within the cell, and uptake by cells. The HPH may include c-raf gene. The HPH molecules function through RNA/DNA inhibition. The HPH molecule binds to and blocks the function of a target nucleic acid, and modulated cellular and viral processes such as splicing, editing, a A method for modulating the function of a target sequence in a cell comprises treatment with a hairpin hybridizer (HPH) nucleic acid molec such as that represented by the present sequence. The present HPH molecule is used in an example of the invention and targets the human sequence that activates RNase H (for cleaving RNA) and greater than for linear antisense sequences. translation. validation, to identify gene function and/or therapeutic targets, The HPH molecule can be used therapeutically, in target quence in a cell nucleic acid molecule HPH may include its specificity editing, and for is

Matches Query Match Best Local 2029 ATCAATGCTTGCACGCTGACCACG 2052 Sequence 31 BP; 6 A; 7 C; 11 G; 1 T; 6 U; 0 other; Similarity Conservative 100.0%; 0.8%; 0, Score 24; Pred. No. Mismatches 3.9e+03 DB 21; 0; Length 31 Indels 0; Gaps

0;

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RESULT 13 AAA29998/c DNA-RNA hybrid; phosphorothicate; hairpin hybridizer molecule; c-raf; inhibitor; detect; nuclease resistance; gene function; gene expression modulator; ss. AAA29998 standard; DNA; 33 misc\_RNA Key Synthetic 09-AUG-2000 Hairpin hybridizer AAA29998 stem\_loop 27 ATCAATGCTTGCACGCTGACCACG (first entry) /note= 1..4 Location/Qualifiers molecule #3 targeting c-raf RNA. "2'-0-methyl-ribonucleotides" ВP HPH;

FTH COS XXX

\*\*\*\*\*\*\*\*\*\*

misc\_feature

/note= 13..21

/\*tag=

/\*tag=

misc\_feature

misc\_feature

/note= 30..33 /\*tag=

'note-

"Phosphorothioate internucleotide linkage"

21-SEP-1998; 20-SEP-1999; 30-MAR-2000 WO200017346-A2

98US-0101174 99WO-US21865 misc\_RNA

/note= 22..33

"Phosphorothioate internucleotide linkage"

"Phosphorothioate internucleotide linkage"

/\*tag=

e "2'-0-methyl-ribonucleotides

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprises treatment with a hairpin hybridizer (HPH) nucleic acid molecule such as that represented by the present sequence. The present HPH molecule is used in an example of the invention and targets the human c-raf gene. The HPH molecules function through RNA/DNA inhibition. The HPH molecule binds to and blocks the function of a target nucleic acid, and modulated cellular and viral processes such as splicing, editing, and translation. The HPH molecule can be used therapeutically, in target validation, to identify gene function and/or therapeutic targets. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     validation, to identify gene function and/or therapeutic targets, for analysis of mutations in diseased cell and to detect specific RNA. The hairpin structure improves resistance to nuclease degradation, localization within the cell, and uptake by cells. The HPH may include a sequence that activates RNase H (for cleaving RNA) and its specificity is
                                                   Thompson JD,
Reynolds M,
                                      Matulic-Adamic
                                                                                         (RIBO-) RIBOZYME PHARM INC
                                                                                                                    20-APR-1998;
23-JUN-1998;
                                                                                                                                                             19-APR-1999;
                                                                                                                                                                                        28-OCT-1999
                                                                                                                                                                                                                   W09954459-A2
                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                         anticancer; breast cancer; endometrium
                                                                                                                                                                                                                                                                                      gene expression modification;
                                                                                                                                                                                                                                                                                                 Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; hammerhead ribozyme; hairpin ribozyme; antisense o
                                                                                                                                                                                                                                                                                                                                       C-raf antisense oligonucleotide sequence SEQ ID
                                                                                                                                                                                                                                                                                                                                                                     19-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                        AAA26239 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2029 ATCAATGCTTGCACGCTGACCACG 2052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A method for modulating the function of a target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 44; Page 76; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modulating target sequence in a c
for identifying gene function, by
hybridizer nucleic acid molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                 Beigelman L,
Zwick M, Jar
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                                                                                                                    98US-0082404
98US-0103636
                                                                                                                                                            .99WO-US08547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear antisense sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                hairpin ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.88;
                                                   Jarvis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INC
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                                                McSwiggen JA,
is T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                      cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment with novel hairpin
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                                                                                                                                                                                                                                                                                    phosphorothicate;
                                                                                                                                                                                                                                                                         cancer;
                                                                                                                                                                                                                                                                                                antisense oligonucleotide;
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                                                Karpeisky A,
Haeberli P;
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                                                                                                                                                                                                                                                                                                                                          NO:2737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ength 33;
                                                                                                                                                                                                                                                                                                               cleavage;
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                                                              Bellon
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RESULT 15
AAA26242/c
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Matches
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                                                                                 Thompson JD, B
Reynolds M, Zw
Matulic-Adamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor. Because of the high selectivity for targeted RNA, (A)
                                                                                                                                                                                             20-APR-1998;
23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C-raf antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2029 ATCAATGCTTGCACGCTGACCAC 2051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and AAA4748 to AAA25992 represent their corresponding target sequences. AAA2593 to AAA2505 represent oestrogen receptor hairpin ribozyme sequences, and AAA26107 to AAA26218 represent their corresponding target sequences. AAA26319 to AAA26271 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that
                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                              anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                           hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonuclec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA26242 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              other ribozyme sequences and antisense oligonucleotides used in exemplification of the present invention.
                                                                                                                                                                                                                                                 19-APR-1999;
                                                                                                                                                                                                                                                                                    28-OCT-1999
                                                                                                                                                                                                                                                                                                                        WO9954459-A2
                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23 BP; 5 A; 4 C; 8 G; 3 T; 3 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes nucleic acids (A) that interact stably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acids that interact, and optionally cleave, target sequences, used to treat cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23
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nes 23; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATCAATGCTTGCACGCTGACCAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 105;
                                                                                     Zwick M,
nic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                      Beigelman L,
                                                                                                                                                                                             98US-0082404
98US-0103636
                                                                                                                                                                                                                                                 99WO-US08547.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide sequence SEQ ID NO:2740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       148pp; English
                                                                                                    n L, McSwiggen JA,
Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%;
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6.6e+03;
                                                                                                    Karpeisky
Haeberli
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                                                                                                                      Bellon L;
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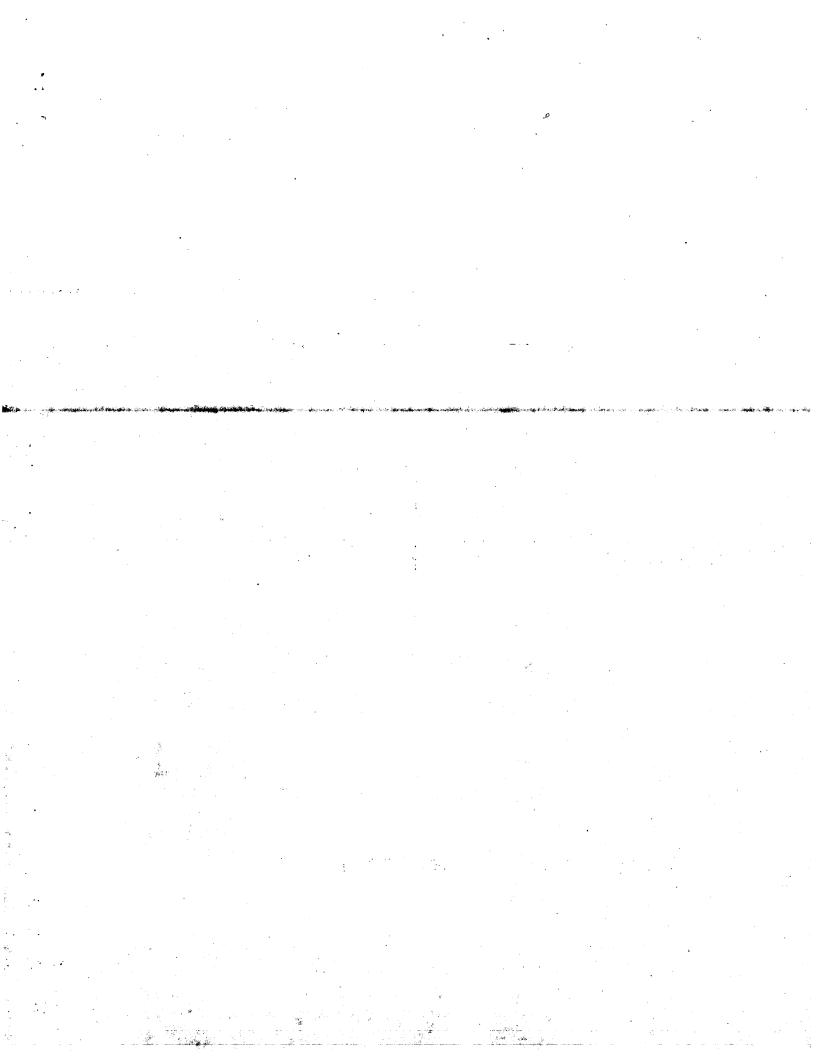
New nucleic acids that interact, sequences, used to treat cancer

and optionally cleave,

target

WPI; 2000-013248/01.

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Search completed: May 16, 2003, 07:08:58 Job time: 644 secs
                                                                                                                                                                                                                                                                                    В
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                                                                                                                                                                                                                                                                                                               The present invention describes nucleic acids (A) that interact stably CC with a target sequence and contain at least one phosphoro(di)thioate CC link, having endonuclease activity. (A), and more generally any CC catalytic nucleic acid (A') that modulates expression of the oestrogen CC receptor gene, are used to treat cancer (particularly of breast or CC endometrium), in vivo or by transforming cells ex vivo and implanting CC treated cells, or for other conditions associated with levels of CC oestrogen receptor. Because of the high selectivity for targeted RNA, (A) CC can also be used to correlate inhibition of gene expression with CC alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that CC restriction endonucleases are used with DNA). The combination of CC modifications in (A) improves resistance to nucleases, binding affinity CA and/Or activity. ANA25503 to ANA24747 represent oestrogen receptor acident ribozyme sequences, and ANA2478 to ANA25992 represent their CC corresponding target sequences. ANA2593 to ANA26107 to ANA26218 represent CC their corresponding target sequences. ANA26219 to ANA26218 represent CC exemplification of the present invention.
                                                                                                                                                                              Query Match 0.8%; Score 23; Best Local Similarity 100.0%; Pred. No. Matches 23; Conservative 0; Mismatc
                                                                                       Sequence 23 BP; 4 A; 8 C; 5 G; 2 T; 4 U; 0 other;
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                                                                                                                                                                                   Mismatches
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                            /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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8612.988 Million cell updates/sec
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Patent No. 6
                               Matches
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                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: ASSAY FOR IDENTIFYING AGENTS WHICH ACT ON TITLE OF INVENTION: CERAMIDE-ACTIVATED PROTEIN KINASE, KINASE
                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957
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                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                            Liu, Jun
Zhang, Yuhua
                                                                                                                                                                                                                                                                                                                                                                                                                              Kolesnick, Richard
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### ALIGNMENTS

SUPPRESSOR OF RAS, AND METHODS OF USING SAID AGENTS

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42 base pairs
42 base pairs
42 base pairs
57RANDEDNESS: single
70POLOGY: linear
70POLECULE TYPE: DNA
US-08-785-247-13
Query M-
Be-
                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/785,247
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1185 Avenue of the Americas
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 Conservative
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Score 35.6; DB Pred. No. 0.099; 0; Mismatches
                          DB 3;
 4; Indels
                            Length 42;
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Gaps
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Best Local Similarity 89.7%;
Matches 35; Conservative
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Patent No. 60
                                                                                                                                                                               GENERAL INFORMATION:
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                                                         CORRESPONDENCE ADDRESS:
                                                                          NUMBER OF SEQUENCES:
                                                                                     TITLE OF INVENTION:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: Cooper & Dunham
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: White, John P. REGISTRATION NUMBER: 28,678 REFERENCE/DOCKET NUMBER: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U
ZIP: 10036
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
              New York
                                                                                                                   INVENTION:
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                                                                                                                                                                                                              Application US/08785247
New York
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                         1185 Avenue of the Americas
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                                                                                                                                  Liu, Jun
Zhang, Yuhua
                                                                                                                                                               Kolesnick, Richard N.
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SYSTEM: PC-DOS/MS-DOS
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                                        Cooper & Dunham LLP
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                                                                                     SUPPRESSOR OF RAS,
                                                                                                ASSAY FOR IDENTIFYING AGENTS WHICH ACT ON THE CERAMIDE-ACTIVATED PROTEIN KINASE, KINASE
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Pred. No. 0.83;
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                                                                                   AND METHODS OF USING
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Best Local Similarity 85.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
   TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
                                                              ATTORNEY/AGENT INFORMATION: NAME: Field, Bret E.
                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                916 GTCCACATGGTCAGCACCACGCTGCCTGTGGACAGCAGGATG 957
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                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
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                                REFERENCE/DOCKET NUMBER:
                                              REGISTRATION NUMBER:
                                                                                                                                          CLASSIFICATION:
                                                                                                                                                     APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997
                                                                                                                                                                                                      OPERATING SYSTEM:
SOFTWARE: FastSE
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EDNESS: single
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2200 Sand Hill Road, Suite 100
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ENTION: METHOD OF AS
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SYSTEM: PC-DOS/MS-DOS
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Pred. No. 1;
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Length 42; Indels

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                                                          OTHER INFORMATION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 979: US-09-225-928-979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 979, Applicat Patent No. 6352829 GENERAL INFORMATION:
               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 979
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1704 CAGTTTCCAGTCGGATGTCTACTCCTAT 1731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CAGTTTCCAGTCGGATGTCTACTCCTAT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: oligonucleotide primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jokhadze, George
Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                               FEATURE:
                                                                                                                           MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                      TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/225,928 FILING DATE: 05-Jan-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                      LENGTH: 28 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                       NAME: Field,
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/859,998 FILING DATE: 21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Windows95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Menlo Park
                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                           ELEFAX: 415-854-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPONDENCE ADDRESS:
NDDRESSEE: Fish & Richardson, P.C.
TREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OF SEQUENCES:
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               Score 28;
Pred. No.
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Pred. No.
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                DB 4;
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RESULT 6
US-08-306-691B-5/c
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                                                                                                                                         Sequence 6, Applic Patent No. 5734039
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                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: NO. 5734039e INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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APPLICANT: Skorski,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: September 15, CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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                                CORRESPONDENCE ADDRESS
                                                               TITLE OF INVENTION:
                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                    26
                                                                                                                                                                                                                                                                                                                                                                                                             TYPE:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: Two Penn Concile Philadelphia STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Monaco, Daniel A. REGISTRATION NUMBER: 30,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                    ATGTGACCGCCTCCCGCTCCCTCACC
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                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
                                                                                                                                                        Application US/08306691B
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                                                                                                                                                                                                                                                                                                                                                                                                                        26 Nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
                                                                                          Skorski, Tomasz
                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                            Calabretta, Bruno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Calabretta, Bruno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WordPerfect 5.1
Seidel, Gonda, Lavorgna & Monaco, P.C.
O Penn Center, Suite 1800
                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                           single stranded
                                                          ANTISENSE OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTISENSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MS-DOS
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                                                                                                                                                                                                                                                                                                                  100.0%;
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                                                                                                                                                                                                                                                                                                                               0.98;
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per 15, 1994
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COUNTRY:

U.S.A.

Pennsylvania

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US-08-859-998-980/
Sequence 980, Application US/08859998
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                                                                                                                                                                                                                                                                                                                                                                                                           ENERAL INFORMATION:
                           ATTORNEY/AGENT INFORMATION:
                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                           SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 26 Nucleotides
                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                             STREET: 2200 Sand Hill Road, Suite 100 STATE: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                              PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ELECOMMUNICATION INFORMATION:
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                                           FILING DATE:
                                                         APPLICATION NUMBER:
                                                                                                APPLICATION NUMBER: US/0
FILING DATE: 21-MAY-1997
                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                         COMPUTER:
                                                                                                                                                                                                                                                                                                                                    TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                     Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          568-5549
                                                                                                                                                         Windows95
                                                                                                                                                                                                                                                                                                                                 METHOD OF ASSAYING DIFFERENTIAL EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MS-DOS
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                                                                                                               us/08/859,998
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                                                                                                                                                                                                                                                                                                                                                             Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 26; ; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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RESULT 9
US-09-225-928-980/c
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Best Local Similarity
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                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 980:
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OTHER INFORMATION: oligonucleotide primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 09096/002001
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                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096,
                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: WI
SOFTWARE: FastSEQ fo
CURRENT APPLICATION DATA:
OTHER INFORMATION: oligonucleotide primer SEQUENCE DESCRIPTION: SEQ ID NO: 980:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                      FEATURE:
                                                         MOLECULE TYPE: DNA
                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid
                                                                                       LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Menlo Park
STATE: CA
                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
                                                                                                                                                                                                    TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
                                                                                                                                                                                     ELEFAX: 415-854-0875
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jokhadze, George
Bibilashvilli, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Windows95
                                                                                                                                                                                                                                                                                                                                                                                                                                             for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 26; DB; Pred. No. 75
0; Mismatches
                                                                                                                                                                                                                                            09096/002001
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75;
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JS-09-482-084-2/c
                                                                                  Sequence 2, Application US/09482084 Patent No. 6333314
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                                                                                                                                                                                                                                      Query Match
Best Local
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Best Local S
                                                                       GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tent No. 6126965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1963 TCTCTACCGAAGATCAACCGGAGCGC 1988
                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                               121 GCTGCATCAATGGAGCACATACAGG 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U
ZIP: 22031
                                                                                                                                                                      25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 24-OCT CLASSIFICATION: 51
                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Hendricks, Glenna REGISTRATION NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                       POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
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                                                        APPLICANT: Kasid, Usha
                                                                        INFORMATION
                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                      (703) 591-4470
(703) 591-4470
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rahman, Aqulur
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                       unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hendricks and Assoc.
             Dritschilo, Anatoly
Rahman, Aqulur
                                        Gokhale, Prafulla
                                                                                                                                                                                                                                                                                                                         DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-OCT-1997
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                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                      0.88;
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Liposomes containng Oligonucelotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           32,535
                                                                                                                                                                                                                                      Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                kasid
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Pred. No.
                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                   Length 25
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                                                                                                                                                                                                                                                                                                                       RESULT 12
US-08-850-049-17
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                                                                                                                                                                                                                                                                            Patent No. 5965726 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                  APPLICANT: APPLICANT:
           SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                         COUNTRY:
                                                                                                                                                   STREET:
APPLICATION NUMBER:
                                                                                                                                                                   ADDRESSEE:
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                               10154
                                                                                                                        NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: P.O. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                   345 PARK AVENUE
                                                                                                            ÜSA
                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                    FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (703)
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US/08/850,049
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Sequence 17, Application US/08850049 Patent No. 5965726
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COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25 GCTGCATCAATGGAGCACATACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIA Release #1.0.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Hendricks, Glenna REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/482,084 FILING DATE: 13-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/957,327 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: kasid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Hendricks and Assoc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                          MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P.O. Box 2509
      PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                      INHIBITORY/INSTABILITY REGIONS OF mRNA
                                                                                                                                                                                                                                                                       METHOD OF ELIMINATING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 591-4428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 25;
Pred. No.
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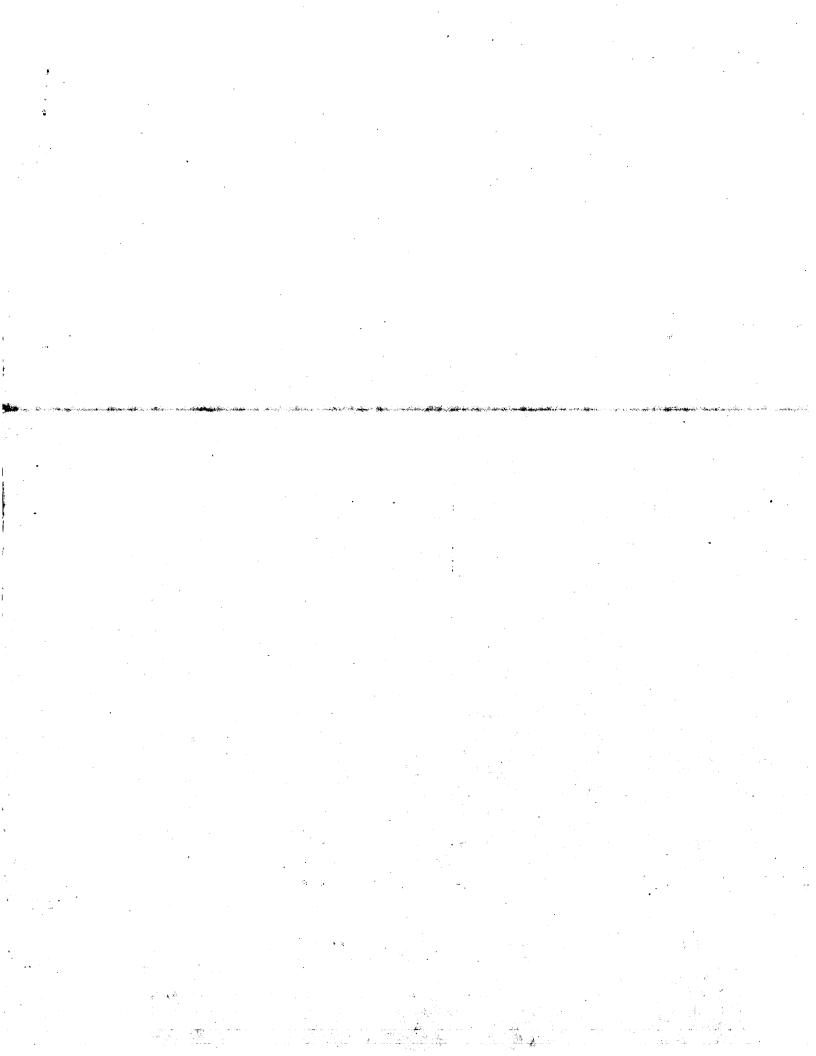
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC 2244
                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 43 BASE PAIRS
                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
TUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT:
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                                              CLASSIFICATION:
                                                                                                                        APPLICATION NUMBER: FILING DATE: 26-OC
                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: MORRY, MARY J.
REGISTRATION NUMBER: 34,398
REFERENCE/DOCKET NUMBER: 20
                PPLICATION NUMBER:
                                                                         PPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 TCTCAGATACCTAGCTTCATATTGCCTTATTGTCTACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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 SSIFICATION DATA:
APPLICATION DATA:
UNMBER: US 07/858,747
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5. 5972596
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                                                                                                                                                                                                                                                                                            NEW YORK
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                                                                                                                                                                                                                                                                                                          345 PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : (212)758-4800
(212)751-6849
OR SEQ ID NO: 1
                                                                                                                                                                     WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LINEAR
                                                                                                                                                                                                                                                                                                                         MORGAN & FINNEGAN
                                                             29-MAR-1993
                                                                                                                      26-OCT-1994
27-MAR-1992
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                                                                                                                                                                                    PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                 INHIBITORY/INSTABILITY REGIONS OF MRNA
                                                                                                                                                                                                                                                                                                                                                                                      METHOD OF ELIMINATING
                                                                         PCT/US93/02908
                                                                                                                                   US/08/050,478
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2026-4006US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 43;
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Best Local Similarity
Matches 28; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6174666
GENERAL INFORMATION:
                                                                                                                                                                                                 NFORMATION FOR
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                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (212)758-4800
                                                                                       HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC 2244
                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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LENGTH: 43 BASE PAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ITLE OF INVENTION:
                                                                                                                                    STRANDEDNESS:
LOCATION:
IDENTIFICATION METHOD:
DTHER INFORMATION: OLIGONUCLEOTIDE FOR MUTATING
                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                        REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                           NAME/KEY:
                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: UFILING DATE: 19920327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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                                                                                                                                                  NUCLEIC ACID
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EDNESS: SINGLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PAVLAKIS, GEORGE N.; FELBER, BARBARA
                                                                                                                                                                                                              (212)751-6849
                                                                                                                    UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                  WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LINEAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FLOPPY DISK
                                                                                                       OTHER
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                                                                                                                                      SINGLE
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FR: 34,398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INHIBITORY/INSTABILITY REGIONS OF mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FINNEGAN
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Pred. No.
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US-09-414-117-17
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                                                                                                                     ;
US-09-414-117-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0. Best Local Similarity 73. Matches 28; Conservative
                                                            Query Match
Best Local Similarity
Matches 28; Conserv
                                                                                                                                                                                                            TELEFAX: (212)751-684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENERAL INFORMATION: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tent No.
2207 TCACAGGGCCTTAACTTCATGTTGCCTTCTTTTCTATC 2244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDPERFECT 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/858,747
FILING DATE: 27-MAR-1992
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/02908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (212)758-4800
                                                                                                                                                                                                                                                                          CLASSIFICATION:
VITORNEY/AGENT INFORMATION:
NAME: MORRY, MARY J.
NAME: MORRY MARY J.
20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                 TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 345 PARK AVENUE
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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CLASSIFICATION:
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                                                                                                                                                                               43 BASE PAIRS
                                                            Conservative
                                                                                                                                                                                                                            (212)751-6849
                                                                                                                                     LINEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08/850,049
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                                                                                                                                                                                                                                                                      2026-4006US1
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Pred. No. 1.9e+03;
                                                                        Score 22; DB 4;
Pred. No. 1.9e+03;
                                                          Mismatches
                                                          10;
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                                                                                        Length 43;
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Search completed: May 16, 2003, 12:18:33 Job time: 113 secs



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Database
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Maximum Match
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Maximum DB
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20.6
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2977
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10213.154 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GenCore version 5:1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_BUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
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0 US-09-263-95-580
0S-09-827-289-19
0 US-09-827-289-23
0 US-09-827-289-23
0 US-09-822-698A-62
US-10-103-906-4
US-10-078-949-5
US-10-078-949-5
US-10-078-949-7
US-09-985-371-37
US-09-986-263-7
US-09-996-263-10
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US-09-930-283A-2
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                                                                                                                                                                                                 Sequence 835, App
Sequence 2, Appli
Sequence 17, Appl
Sequence 213, App
Sequence 10177, A
Sequence 580, App
                                                          Sequence 4, Appli
Sequence 3, Appli
Sequence 5, Appli
Sequence 7, Appli
Sequence 37, Appl
            Sequence
Sequence
Sequence
Sequence
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                                                                                                                                       Sequence 99, Appl
Sequence 19, Appl
Sequence 23, Appl
Sequence 62, Appl
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           37, Appli
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US-09-930-283A-2/c
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US-09-801-274-835
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LENGTH: 31
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Best Local :
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ALIGNMENTS	10-057	10-057	057	)-057	057	057	057	057	057	057	10-057	10-057	10-057	0-057	10-057	0-057	0-057	0-057	0-057	932	071	835	996	996	96	996
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# ALIGNMENTS

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; ORGANISM: Homo sapiens US-09-801-274-835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
APPLICANT: Lander, HUMAN SINGLE NUCLEOTIDE
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE
FILE REFERENCE: 2825, 2009-001
                                                                                   Sequence 2, Application US Patent No. US20020160038A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 835, Application US/09801274 Patent No. US20020032319A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DAYE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                          2144 CCCTTTCTCCAGAGGCAGAACACATGTTTTC 2174
                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                   Conservative
: Kasid, Usha
Gokhale, Prafulla
Dritschilo, Anatoly
Rahman, Aqulur
                                                                                                                                                                                                                                                                                                                                    1.0%;
                                                                                                                            US/09930283A
                                                                                                                                                                                                                                                                                                              Score 30.6; D
Pred. No. 13;
1; Mismatches
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US-09-943-722-17
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION:
                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                        PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 GCTGCATCAATGGAGCACATACAGG 145
                                OPERATING SYSTEM:
                                                   COMPUTER:
                                                                                                                   COUNTRY:
                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                 25 GCTGCATCAATGGAGCACATACAGG 1
                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL:
ANTI-SENSE: Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 1999-07-15
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Hendricks and Assoc
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ARE: WORDPERFECT 5.1
APPLICATION DATA:
                                                                                                    10154
                                                                                                                                   NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: ka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/930,283A FILING DATE: 16-Aug-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (703)
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                                                                                                                                                                                                                                                                                                                                  Application US/09943722
No. US20020192660A1
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                                                                                                                                                                  345 PARK AVENUE
                                                                                                                 USA
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                                                IBM PC COMPATIBLE
                                                                                                                                                                                 MORGAN & FINNEGAN
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                                PC-DOS/MS-DOS
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                                                                                                                                                                                                                METHOD OF ELIMINATING INHIBITORY/INSTABILITY REGIONS OF mRNA 130
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Pred. No.
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7.1e+
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US-09-765-081-213
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                                                                                           Matches
                                                                                                                          Query Match
                                                                                                                                                                                                                           SEQ ID NO 213
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                                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/765,081
CURRENT FILING DATE: 2001-01-18
PRIOR APPLICATION NUMBER: US 60/176,861
PRIOR FILING DATE: 2000-01-19
                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. US20020037508A1
                                                                                                                                                                                          LENGTH: 21
TYPE: DNA
                                                                                                                                                                                                                                          NUMBER OF SEQ
SOFTWARE: Fast
                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ireland, James S.
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2008-001
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cargill, Michele
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                         ORGANISM: Homo sapiens
                                           1237 GGAGATGTTGCAGTAAAGATC 1257
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LENGTH: 43 BASE PAIRS
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APPLICATION NUMBER:
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TYPE: N
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REGISTRATION NUMBER: 34,398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                         Conservative
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                                                                                                                                                                                                                                          for Windows Version 4.0
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                                                                                                         95.2%;
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                                                                                                         Score 20.6;
Pred. No. 1
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Pred. No. 9.
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3e+03

Length 43;

10;

Indels

0;

0;

6e+04

Indels Length

Gaps.

DB 10;

21; 0;

Matches

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US-09-263-959-580/c
                                                               US-09-263-959-580
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US-09-764-891-10177
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              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prior application data removed - consult PALM NUMBER OF SEQ ID NOS: 10231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/764,891
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TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PC006
                                                                                                                                                         NFORMATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . 2868 ACAGGTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATTG 2910
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry LLP
                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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les 29; Conserv
                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                 TORNEY/AGENT
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 05-MAI
                                                                               TOPOLOGY:
                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 92
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                                                                                                                                                                                      TELEPHONE:
                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                           LENGTH:
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                                                                                                           nucleic acid
                                                                                                                                                                                                                                               McMasters, David D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    I: Rowen, Lee T: Koop, Ben F. INVENTION: DIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington
                                                                                                                        31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6300 Columbia Center,
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                                                                              linear
                                                                                                                                                          SEQ ID NO:
                                                                                                                                                                                                                                                                 INFORMATION:
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                                                                                           single
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                0.7%;
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              Score 20.4;
Pred. No. 2.
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Pred. No. 2
   Mismatches
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                4e+04
                              DB IO;
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OTHER INFORMATION:
OTHER INFORMATION:
US-09-827-289-19
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US-10-085-906-99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-827-289-19
                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. SEQ ID NO 19
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                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. US20020009716A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No.
                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/827,289
CURRENT FILING DATE: 2001-04-05
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Abarzua, Patricio
TITLE OF INVENTION: Process for Allele Discrimination Using
TITLE OF INVENTION: Extension
FILE REFERENCE: 469290-55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ying, Vin
                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-0
                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: U.S. 60/194843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-03-24
                                                                                                                                                                                               LENGTH: 46
TYPE: DNA
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                                                                                                                                                            FEATURE:
                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIOR FILING DATE: 1999-03-25
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2893 TTAATTTTGTTTTTATTGCACCTGACAAAATACAGTTATCT 2933
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US00/07938
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                                                    Similarity
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FastSEQ for Windows Version 4.0
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                                                                                                                        Description of Artificial Sequence: Pl primer for use in allele discrimination
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                                                   Score 20.2; DB 10;
Pred. No. 3.7e+04;
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Pred. No. 2
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RESULT 11
US-10-103-906-4/c
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US-09-822-698A-62
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                                                       Sequence 4, Application US/10103906
Patent No. US20020156268A1
                                              GENERAL
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Best Local Similarity
Matches 28; Conser
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APPLICANT: Abarzua, Patricio
TITLE OF INVENTION: Process for Allele Discrimination Using Primer
TITLE OF INVENTION: Extension
                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. US20020146750A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 62,
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Sequence 23, Application U
Patent No. US20020009716A1
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PRIOR FILING DATE: 2000-03-30
NUMBER OF SEQ ID NOS: 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: MUCIN-1 SFILE REFERENCE: DYX-015.1 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Hoogenboom, Hendricus R.J.M. APPLICANT: Henderikx, Maria P.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: nucleotide sequence coding
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                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 48
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                                           INFORMATION
                                                                                                                                                           GGGGGGGCCAGAAGCTGCAGCTGGACTACTGG
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Krotz, Achim н.
McElroy, Bethany
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                                                                                                                                                                                                                               Conservative
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Pred. No. 3
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RESULT 13
US-10-078-949-5
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                                                                                                               GENERAL
                                                                                                                            Sequence 5, Application US/10078949 Patent No. US20020165189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 3
LENGTH: 20
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SEQ ID NO 4
LENGTH: 20
CURRENT APPLICATION NUMBER: US/10/078,949
CURRENT FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: 09/479,783
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                                                      TITLE OF INVENTION: Oligoribonucleotides And Ribonucleases For Cleaving RNA FILE REFERENCE: ISIS5027
                                                                                       APPLICANT: Crooke, Stanley T.
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PRIOR FILING DATE: 1997-06-06
PRIOR APPLICATION NUMBER: 08/659,440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Crooke, Stanley T.
TITLE OF INVENTION: Oligoribonucleotides and Ribonucleases For Cleaving RNA
FILE REFERENCE: ISIS5027
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CURRENT FILING DATE: 2002-02-20
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                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
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CURRENT FILING DATE: 2002-03-22
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nes 20; Conserv
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                                                                                                            INFORMATION:
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                                                                                                                                                             US-09-835-371-37/c
; Sequence 37, Application US/09835371
; Publication No. US20020187473A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Oligonucleotide US-10-078-949-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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Patent No. US20020165189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1 SEQ ID NO 5
                              APPLICANT: UHLMANN, Eugen
APPLICANT: BREIPOHL, Gerhard
APPLICANT: WILL, David W
APPLICANT: WILL, David W
TITLE OF INVENTION: POLYAMIDE NUCLEIC ACID DERIVATIVES,
TITLE OF INVENTION: PROCESSES FOR PREPARING THEM
FILE REFERENCE: 02481.1743 SEQUENCE LISTING
CURRENT APPLICATION NUMBER: US/09/835,371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -10-078-949-7
CURRENT FILING DATE: 20 NUMBER OF SEQ ID NOS: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 08/870,608
PRIOR FILING DATE: 1997-06-06
PRIOR APPLICATION NUMBER: 08/659,440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/078,949
CURRENT FILING DATE: 2002-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/479,783 PRIOR FILING DATE: 2000-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Crooke, Stanley T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
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FILING DATE: 1997-06-06
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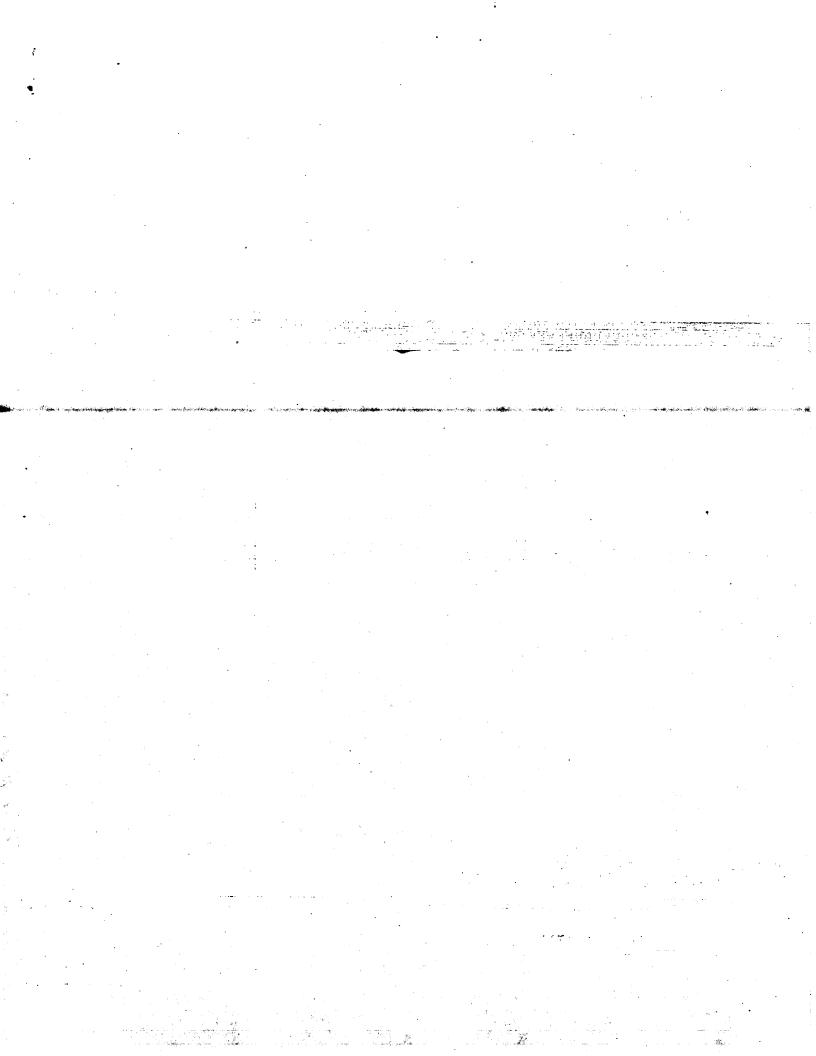
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LENGTH: 20
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                                                               Local Similarity
                                                     Conservative
                                                                 100.0%;
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Search completed: May 16, 2003, 12:48:11 Job time: 381 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result

Query

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                                                                                                                                                            TTGTTTTTATTGCACCTGACAAAATACAGTTATCTGATGGTCCCTCAATT 2948
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; SEQ ID NO 3119

; LENGTH: 50

; TYPE: DNA

; ORGANISM: Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Homo US-10-131-827-3119
GENERAL INFORMATION:
APPLICANT: Glenn, Mattew
APPLICANT: Grigor, Murray R
TITLE OF INVENTION: Compositions isolated from bovine tissue
TITLE OF INVENTION: and methods for their use.
FILE REFERENCE: 1055P3
                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 50
                                                                                                                Sequence 9462, Application US/60253651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1
SEQ ID NO 3119
LENGTH: 50
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Wohlgemuth, Jay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 10/006,290 PRIOR FILING DATE: 2001-10-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ly, NGOC
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: APPLICANT:
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APPLICANT: Ery, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
TITLE OF INVENTION: CHRONIC INFLAMMATORY DISEASES
FILE REFERENCE: 506612000120
CURRENT APPLICATION NUMBER: 05/10/131,827
CURRENT FILING DATE: 2002-09-06
CURRENT FILING DATE: 2002-09-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 506612000121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-66-08
NUMBER OF SEQ ID NOS: 9990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wohlgemuth, Jay
                                                                                                                                                                                                                                                        2899 TTGTTTTTATTGCACCTGACAAAATACAGTTATCTGATGGTCCCTCAATT 2948
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                                                                                                                                                                                                                50 TIGITTTTATIGCACCTGACAAATACAGTTATCTGATGGTCCCTCAATT 1
                                                                                                                                                                                                                                                                                                Local Similarity 100. 
les 50; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    50 TTGTTTTTATTGCACCTGACAAAATACAGTTATCTGATGGTCCCTCAATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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Woodward, Robert
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                                                                                                                                                                                                                                                                                                                DB 40;
0.017;
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0.017;
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RESULT 1
PCT-US01-47856-3119/c
Sequence 3119, Application PC/TUS0147856

ALIGNMENTS

APPLICANT: BIOCARDIA,

INC

Wohlgemuth, Jay Quertermous, Thomas Johnson, Frances

RESULT 2
US-10-131-827-3119/c
; Sequence 3119, Appli;
; GENERAL INFORMATION:

Application US/10131827

밁 Qy

2899 50

; TYPE: DNA; Homo sapiens PCT-US01-47856-3119

SOFTWARE: PatentIn version 3.1 EQ ID NO 3119 LENGTH: 50

CURRENT APPLICATION NUMBER: PCT/US01/47856
CURRENT FILING DATE: .2001-10-22
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER: OF SEQ ID NOS: 8832

APPLICANT:

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Fry,

Matcuk, George Prentice, James Phillips, Julie Woodward, Robert

APPLICANT: Ly, Ngoc
APPLICANT: Altman, Peter
TITLE OF INVENTION: LEUKOCYTE EXPRESSION PROFILING
FILE REFERENCE: 506612000140

Best Local Similarity Matches 50; Conserv

Conservative

1.7%; Score 50; 100.0%; Pred. No. 0;

DB 1; 0.017;

Mismatches

0,

Query Match

CURRENT APPLICATION NUMBER: US/60/253,651 CURRENT FILING DATE: 2000-11-28 NUMBER OF SEQ ID NOS: 27858 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 9462

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; ORGANISM: Bovine
US-60-253-651-9462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-801-274-835
                                                                                                                            NUMBER OF SEQ ID NOS: 30124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25508
                                Query Match
                                                                                                                                                                                                                                                        Sequence 25508, Application GENERAL INFORMATION:
APPLICANT: Glenn, Matthew
                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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      Matches
                                                                 ORGANISM: Ovine -60-253-653-25508
                                                                                                                                                                                                                                                                                                       60-253-653-25508
                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/60/253,653
CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/206,129 PRIOR FILING DATE: 2000-05-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/187,510 PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                                             FILE REFERENCE: 1057P
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                                                                                                 LENGTH: 46
TYPE: DNA
                                                                                                                                                                                                                         TITLE OF INVENTION: Compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PPLICANT: Ireland, James S.
PPLICANT: Lander, Eric S.
ITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
ILE REFERENCE: 2825.2009-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 48
                                                                                                                                                                                                                                                                                                                                                                                                    2144 CCCTTTCTCCAGAGGCAGAACACATGTTTTC 2174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity hes 37; Conserv
                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                      CCCTTTCTCCAGAGGYAGAACACATGTTTTC 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGACCCGAACCCCTACAGCTTCCAGTCAGATGTCTACGCCTATGG 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/801,274 FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FastSEQ for Windows Version 4.0
    Conservative
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                   Score 30.6;
Pred. No. 2
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Pred. No. 2
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    Mismatches
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                   .8e+03
                                    DB 69;
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                                  Length 46;
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US-09-225-201B-979
; Sequence 979, Application US/09225201B
; GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
                                                                                                                          US-08-306-691-5/c; Sequence 5, Application US/08306691; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: C;
SEQUENCE DESCRIPTION: SEQ
US-09-225-201B-979
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                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        :NFORMATION FOR SEQ ID NO: 979:
                                                        TITLE OF INVENTION:
                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                             1 CAGTTTCCAGTCGGATGTCTACTCCTAT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
ZIP: 94025
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCACTCCCATTGGATCACGCTCCTTCGGGACTGTTTATAACGGCA
                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/225,201B FILING DATE: 05-Jan-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Field,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 28 base pairs
                                                                                       Skorski,
                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                               Calabretta, Bruno
                                                                                                                                                                                                                                                                                                               0.9%; 500
100:0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
                                                        ANTISENSE .
OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
Gonda, Lavorgna & Monaco, P.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bret E.
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                                                                                                                                                                                                                                                                                                                                       Score 28; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide primer
Q ID NO: 979:
                                                                                                                                                                                                                                                                                                                     Mismatches
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COMPUTER READABLE FORM:

: Two Penn Center, Suite 1800 Philadelphia

Pennsylvania

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US-08-306-691-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 Nucleans
                                                                                                                                                                                                                                                                                                                                                                                              quence 6, Applicati
ENERAL INFORMATION:
                        ATTORNEY/AGENT INFORMATION:
                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                         ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NTTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       ITLE OF INVENTION:
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                                                                                                                                                             OPERATING SYSTEM:
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SOFTWARE: Wordpor
                                                         APPLICATION NUMBER:
                                                                                                     FILING DATE:
                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Monaco, Daniel REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IOR APPLICATION DATA: APPLICATION NUMBER:
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                                                                                                                                                                                                                       19102
                                                                                                                                                                                                                                                           SEE: Seidel, Gonda,
: Two Penn Center,
Philadelphia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08306691
                                                                                                                                                                                                                                                 Pennsylvania
                                                                                                                                                                                                                                                                                                                        SEQUENCES:
                                                                                                                                                                                                                                  U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (215) 568-5549
                                                                                                                                             WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                                               Calabretta, Bruno
                                                                                                                                                                        IBM PS/2
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IBM PS/2
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                                                                                                                                                             MS-DOS
                                                                                                                                                                                                                                                                                                                                                     ANTISENSE
                                                                                                                                                                                                                                                                                                                                   OLIGONUCLEOTIDES TARGETING COOPERATING ONCOGENES
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                           Gonda,
                                                                                                                 US/08/306,691
A.
30,480
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                                                                                                                                                                                                                                                                            Suite 1800
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  Matches
                             Query Match
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GENERAL INFORMATION
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                                                                                                                                            INFORMATION FOR SEQ ID NO:
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                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-8383
                                                                                                                                                                                                                                                      FILING DATE: September 15, 1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
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LENGTH: 26 Nucleotides
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                                                                                    STRANDEDNESS:
                Local Similarity
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                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PS/2
OPERATING SYSTEM:
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                                                                                                                                                        None
                                                                                                                                                                                                                                             Monaco,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pennsylvania
                                                                                                              26 Nucleotides
                                                                                                                                                                      (215) 568-5549
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                                                                        linear
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                                                                                                                                                                                                                    Daniel A. 30,480
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              0.9%;
100.0%;
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             Score 26;
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Pred. No.
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 Mismatches
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                             DB 12;
                3.6e+04
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                                                                                                                                                                                                  Sequence 980, Application US/09225201B
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                    09-225-201в-980/с
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ENERAL INFORMATION:
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COMPUTER: 15..
OPERATING SYSTEM: MS-L.
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                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish &
COMPUTER READABLE FORM:
                                                                                                                           NUMBER OF SEQUENCES: 1375
                                                                                                                                                    Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAVING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                  ATGTGACCGCCTCCCGCTCCCTCACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
              COUNTRY: US
ZIP: 94025
                                                             CITY: Menlo Park
                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         None
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                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                     Jokhadze, George
Bibilashvilli, Ro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    September 15,
                                                                          EE: Fish & Richardson, P.C.
2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single stranded
                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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PROLIFERATION WITH ANTISENSE OLIGONUCLECTIDES TARGETING COOPER
                                                                                                                                                                                                                                                                                                                                                                                                               0.98;
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                                                                                                                                        EXPRESSION
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Pred. No.
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Suite 1800
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                                                                                                                        CLASSIFICATION:
                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                               STREET: P.O. CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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Sequence 2, Application:
                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/957,327
FILING DATE: 24-OCT-1997
ATTORNEY/ACENT INFORMATION:
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                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ELECOMMUNICATION INFORMATION:
             REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                     APPLICATION NUMBER:
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FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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SEQUENCE DESCRIPTION: SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09354109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Field, מוכנ ב.
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
                                                    Hendricks, Glenna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 05-Jan-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/225,201B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rahman, Agulur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dritschilo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gokhale, Prafulla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kasid, Usha
                                                                                                                                                                                                                                                           IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                             Hendricks and Assoc
                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                          PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Liposomes containng Oligonucelotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%; Score 26;
100.0%; Pred. No.
                                                                                                                                                                                     US/09/354,109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anatoly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide primer
Q ID NO: 980:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                          Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Gaps

0

TELEFAX: (703) 591-4428 INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

ENGTH:

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RESULT 15
US-09-396-196F-44995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; NUMBER OF SEQ ID NOS: 127806
; SOFTMARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 44994
; LENGTH: 25
TYPE: DNA
; ORGANISM: mus musculus
US-09-396-196F-44994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Qγ
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US-09-354-109-2
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0%;
Matches 25; Conservative
                                                                                                                                                                                                                                                             Sequence 44995, Application US/09396196F GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 44994, Application US/09396196F GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8
Best Local Similarity 100.
Matches 25; Conservative
                                                SEQ ID NO 44995
                                                                             PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
                                                              SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                         FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196F
CURRENT FILING DATE: 2001-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/396,196F CURRENT FILING DATE: 2001-09-15
                                                                                                                                                                         APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
AFFINE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                             APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/100,678 PRIOR FILING DATE: 1998-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
ORGANISM: mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Michael Mittmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NPPLICANT: Affymetrix, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: un
MOLECULE TYPE:
HYPOTHETICAL: 1
                                                                                                                                                                                                                                                                                                                                                                            1855 AACTGCCCCAAAGCAATGAAGAGGC 1879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121 GCTGCATCAATGGAGCACATACAGG 145
                                                                                                                                                                                                                                                                                                                                                           1 AACTGCCCCAAAGCAATGAAGAGGC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25 GCTGCATCAATGGAGCACATACAGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    H: 25 base pairs nucleic acid NDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  David Mack
David Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             YES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Jr
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; ; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DВ 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 17;
6.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                           6.5e+()4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                        ,Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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US-09-396-196F-44995
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Query Match 0.8%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 6. Matches 25; Conservative 0; Mismatches DB 17; 1 Length 25; 0;

Gaps

0;

1876 AGGCTGGTAGCTGACTGTGTGAAGA 1900

B 1 AGGCTGGTAGCTGACTGTGAAGA 25

Search completed: May 16, 2003, 12:16:27 Job time : 6348 secs

GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on: May 16, 2003, 07:11:11; Search time 1366 Seconds (without alignments) 11277.539 Million cell updates/sec

US-10-057-550A-64 2977

Title: Perfect score: Sequence:

Scoring table: IDENTITY\_NUC
Gapop 10.0 , Gapext 1.0

Searched: 6404235 seqs, 2587356060 residues

Total number of hits satisfying chosen parameters: 7758442

Minimum DB seq length: 0 Maximum DB seq length: 50

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Pending\_Patents\_NA\_New:\* /cgn2\_6/ptodata/2/pna/PCT\_NEW\_COMB.seq:\*
/cgn2\_6/ptodata/2/pna/USG6\_NEW\_COMB.seq:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq2:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq3:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq3:\*
/cgn2\_6/ptodata/2/pna/USG9\_NEW\_COMB.seq3:\*

/ptodata/2/pna/US10\_NEW\_COMB.seq2:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result		Score	% Query Match	Length	DВ	ID	Description
n	ш	50	1.7	50	9	US-10-325-899-3119	Sequence 3119, Ap
C	2	25	0.8	25	80	US-10-075-994A-2	2, Apı
	ω	23.4	0.8	25	11	US-60-427-808-320641	Sequence 320641,
	4	23.4	0.8	25	11	US-60-427-808-556623	
	J	23.4	0.8	25	11	US-60-427-808-917222	
	6	23.4	0.8	25	11	US-60-427-836-397923	Sequence 397923,
ი		23	0.8	50	9	US-10-325-899-5803	Sequence 5803, Ap
		22.4	0.8	43	.11	US-60-434-832-1070	Sequence 1070, Ap
	_	22.2	0.7	45	=	US-60-288-292-19321	Sequence 19321, A
		22	0.7	22	9	US-10-293-338-4068	Sequence 4068, Ap
	Ξ	22	0.7	47	9	US-10-349-143-3876	Sequence 3876, Ap
	12	21.8	0.7	25	_	PCT-US02-14877A-875	Sequence 875, App
		21.8	0.7	25	11	US-60-427-808-320640	Sequence 320640,
	14	21.8	0.7	25	11	US-60-427-808-556622	Sequence 556622,
	5	21.8	0.7	25	11	US-60-427-808-917221	Sequence 917221,
	16	21.8	0.7	25	11	US-60-427-836-397922	Sequence 397922,
	17	21.8	0.7	39	æ	US-10-287-787-18691	Sequence 18691, A
0	8		0.7	39	œ	US-10-287-787-18692	Sequence 18692, A
	19	21.8	0.7	. 39	æ	US-10-287-787-18693	Sequence 18693, A
	20	21.8	0.7	50	œ	US-10-287-787-19984	Sequence 19984, A
o 	21	21.6	0.7	50	σ	US-09-912-293-210904	Sequence 210904,
	22	21.6	0.7	50	9	US-10-325-899-4714	Sequence 4714, Ap

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45	44	43	42	41	40	39	38	37	36	35	34	<b>ω</b> <b>ω</b>	32	31	30	29	28	27	26	25	24	23
20.4	20.4	20.4	20.4	20.6	20.6	20.6	20.8	20.8	20.8	20.8	20.8	21	21	21	21	21	21	21.2	21.2	21.2	21.4	21.4
0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7	0.7
50	50	45	45	50	50	<u>4</u> 5	50	25	25	25	25	50	50	50	47	44	44	49	47	46	50	25
9	9	8	8	9	6	8	9	11	11	11	11	9	9	σ	9	8	œ	8	8	æ	0	11
US-10-325-899-4603	US-10-325-899-3367	US-10-287-787-21970	US-10-287-787-21968	US-10-325-899-1696	US-09-912-293-43884	US-10-287-787-19197	US-10-325-899-7636	US-60-427-836-451365	US-60-427-808-943561	US-60-427-808-630023	US-60-427-808-506441	US-10-325-899-8045	US-10-325-899-3101	US-09-912-293-242736	US-10-349-143-2948	US-10-287-787-19790	US-10-287-787-19789	US-10-389-048-4355	US-10-287-787-4936	US-10-287-787-4937	US-09-912-293-130705 ·	US-60-427-808-116035
Sequence 4603, Ap	Sequence 3367, Ap	Sequence 21970, A	Sequence 21968, A	Sequence 1696, Ap	Sequence 43884, A	Sequence 19197, A	Sequence 7636, Ap	Sequence 451365,	Sequence 943561,	Sequence 630023,	Sequence 506441,	Sequence 8045, Ap	Sequence 3101, Ap	Sequence 242736,	Sequence 2948, Ap	Sequence 19790, A	Sequence 19789, A	Sequence 4355, Ap	Sequence 4936, Ap	Sequence 4937, Ap	Sequence 130705,	Sequence 116035,

## ALIGNMENTS

•	•				
RESULT 2 US-10-075-994A-2/c Sequence 2, Application US/10075994A GENERAL INFORMATION: APPLICANT: GOKHALE, Prafulla APPLICANT: GOKHALE, Prafulla APPLICANT: MEMANI, Rajshree APPLICANT: AHMAD, Imran	Hest Local Similarity 100.0%; Freq. No. 0.00051; Matches 50; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 2899 TTGTTTTATTGCACCTGACAAATACAGTTATCTGATGGTCCCTCAATT 2948	TYPE: DA  ORGANISM: Homo sapiens US-10-325-899-3119  Ouery Match  1.7%; Score 50; DB 9; Length 50;	; PRIOR APPLICATION NUMBER: US 10/006,290 ; PRIOR FILING DATE: 2001-10-22 ; PRIOR APPLICATION NUMBER: US 10/131,831 ; PRIOR FILING DATE: 2002-04-24 ; NUMBER OF SEQ ID NOS: 9966 ; SOFTWARE: PATENTIN VERSION 3.1 ; SEQ ID NO 3119	OF INVENTION: REJECTION REFERENCE: 506612000122 NT APPLICATION NUMBER: US/10/325,899 NT FILING DATE: 2002-12-20 APPLICATION NUMBER: US 60/296,764 FILING DATE: 2001-06-08	RESULT 1  US-10-325-899-3119/c  IS SEQUENCE 3119, Application US/10325899  GENERAL INFORMATION:  APPLICANT: Wohldemuth, Jay  APPLICANT: Fry, Kirk  APPLICANT: Ly, Ngoc  APPLICANT: Ly, Ngoc  APPLICANT: Woodward, Robert  TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPI.

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; ORGANISM: Mus musculus US-60-427-808-556623
                                                                                                                                                                                                                                                                                                                                                                     밁
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US-60-427-808-320641
                                                                                                                                                                                                                                                                                                        US-60-427-808-556623
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                                             Query Match
Best Local :
                               Matches
                                                                                                                                                      NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe S
SEQ ID NO 556623
                                                                                                                                                                                                                                                              Sequence 556623, Ap GENERAL INFORMATION APPLICANT: Xue Mei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Microarray Probe SEQ ID NO 320641
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 320641,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.2
SEQ ID NO 2
LENGTH: 25
                                                                                                                        LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                          TITLE OF INVENTION: Methods of Genetic Analysis of Mouse FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 982914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Methods of Genetic Analysis FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xue Mei Zhou
2214 GCCTTAACTTCATGTTGCCTTCTTT 2238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: RAHMAN, AGUIIUT
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
FILE REFERENCE: 219604
                                                                                                                                                                                                                                                                                                                                                                               2210 CAGGGCCTTAACTTCATGTTGCCTT 2234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121 GCTGCATCAATGGAGCACATACAGG 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25
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                                                                                                                                         25
                                            Similarity
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                                                                                                                                                                                                                                                             Xue Mei Zhou
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                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                      Application US/60427808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/60427808
                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%;
                                            0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%;
                                                                                                                                                                 Sequence Listing Generator V 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence Listing Generator V 1.1
                              0;
                                            Score 23.4;
Pred. No. 2.
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Pred. No. 2
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Pred. No.
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                                          2.4e+04;
                                                          DB 11;
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; ORGANISM: Rattus norvegicus
US-60-427-836-397923
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GENERAL INFORMATION
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Best Local Similarity
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LENGTH: 25
                                                 FILE REFERENCE: 506612000122
CURRENT APPLICATION NUMBER: US/10/325,89:
CURRENT FILING DATE: 2002-12-20
                                                                                              APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPL
TITLE OF INVENTION: REJECTION
                     PRIOR APPLICATION NUMBER: US 60/296,764 PRIOR FILING DATE: 2001-06-08
                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/60/427,836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methods of Genetic Analysis of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
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TITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                              2214 GCCTTAACTTCATGTTGCCTTCTTT 2238
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APPLICATION NUMBER: US 10/006,290
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Pred. No. 2
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SOFTWARE: PatentIn version 3.1 SEQ ID NO 5803

LENGTH: 50

NUMBER OF SEQ ID NOS: 9966

PRIOR FILING DATE:

PRIOR APPLICATION NUMBER: US 10/131,831

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TCGCCCACCGAGATATTAAATCCAGCCACATAATGCTCTTTGA 45

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NUMBER OF SEQ ID NOS: 7137

SOFTWARE: PatentIn version 3.1
SEQ ID NO 1070
LENGTH: 43
TYPE: DNA
ORGANISM: Candida albicans
US-60-434-832-1070
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US-60-288-292-19321
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Best Local S
Matches 29
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                                                                                                                                                   SEQ ID NO 19321
LENGTH: 45
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Best Local Similarity
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                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/288,292
CURRENT FILING DATE: 2001-05-02
                                                                                                                                                                                                                                         TITLE OF INVENTION: Compounds TITLE OF INVENTION: and metherile Reference: 1058P
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                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                                                                                                                                                                                                                                                                   APPLICANT: Glenn, Matthew APPLICANT: Norriss, Michael G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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1520 TCATCCATAGAGACATGAAATCCAACAATATATTTCTCCCATGA 1562
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                                                              Local Similarity
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                                              Conservative
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Pred. No.
                                                            Score 22.2; DB 11
Pred. No. 7.1e+04;
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Pred. No. 6.1e+04;
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PCT-US02-14877A-875; Sequence 875, Applic; GENERAL INFORMATION:

Application PC/TUS0214877A

RESULT 12

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US-10-293-338-4068
                                                                                                                                              ; OTHER INFORMATION: 99-4254-307 : US-10-349-143-3876
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US-10-293-338-4068
                                                                                                                                                                                                                                                                            NUMBER OF SEQ
SEQ ID NO 3876
LENGTH: 47
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CURRENT FILING DATE: 2007-11-14
NUMBER OF SEQ ID NOS: 8785
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4068
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-23
PRIOR FILING DATE: EARLIER FILING DATE: 1998-11-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/349,143 CURRENT FILING DATE: 2003-01-21
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TITLE OF INVENTION: THEREOF
                                                                                                                                                                               FEATURE:
NAME/KEY: allele
LOCATION: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: GENSET.020CP1
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TYPE: DNA
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                                                                                                                                                                                                                                         ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ITLE OF INVENTION: Biallelic markers for use
                                2554 AACAGCCCAGGACTGCCGAGACTCTGGCCGCCGAAGGAGCCTGCT 2599
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AACAGCCTTGGGCTCACGTTACARCAGCATCCCCAAGGAGCCTCCT 46
                                                                                        Similarity
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Pred. No.
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Pred. No. 8.3e+04;
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1998-04-21
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; TYPE: DNA
; ORGANISM: Mus musculus
US-60-427-808-320640
                                                                                                                                                                                                                                                       US-60-427-808-556622
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                                                                                                                                                                                  Sequence 556622, Application US/60427808 GENERAL INFORMATION:
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SOFTWARE: Microarray Probe
SEQ ID NO 320640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
CURRENT APPLICATION NUMBER: US/60/427,808 CURRENT FILING DATE: 2002-11-20
                                                                        APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 320640, Application US/60427808 GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/427,808
CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anisowicz, Anthony
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Search completed: May
Job time : 1374 secs
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US-60-427-808-917221
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US-60-427-808-917221
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 556622
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Matches 23; Conservative
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                                                                                                                                                                                                                       SOFTWARE: Microarray Probe Sequence Listing Generator V.1.1 \pm Q ID NO 917221
                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                 TYPE: DNA ORGANISM:
                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 98291
                                                                                                                                                                                                                                                                                              FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Mus musculus
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                                                                               2217 TTAACTTCATGTTGCCTTCTTTTCT 2241
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AZ391073 1M0152I24
BE969098 601650077
BJ047667 BJ047667
                                                                                                                                                                                                    Description
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AUTHORS
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KEYWORDS
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DEFINITION
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JOURNAL
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Fax: 314 286 1810
Email: est@watson.wustl.edu
Insert Size: 603
High quality sequence starts: 1
                                                                                                                                                                                                                                                                          Mammalia; Butheria; Primates; Catarrhin; Hominidae; Homo.

1 (bases 1 to 48)

2 (bases 1 to 48)

2 (bases 1 to 48)

3 (bases 1 to 48)

4 (bases 1 to 48)

5 (bases 1 to 48)

6 (bases 1 to 48)

7 (bases 1 to 48)

7 (bases 1 to 48)

8 (bases 1 to 48)

1 (bases 1 
                                                                                                                                                                    Unpublished (1995)
Contact: Wilson RK
Washington University School of Medicine
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      H26959
H26959.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     48 bp mRNA linear EST J yo58c08.sl Soares breast 3NbHBst Homo sapiens cDNA clone IMAGE:182126 3' similar to gb:X06409 RAF PROTO-ONCOGENE - SERINE/THREONINE-PROTEIN KINASE (HUMAN);, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AA139046
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AA853120
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AU2770282 AU2770282
AZ310534 TM0025503
AL587842 AL587842
AZ512567 1M0358617
A1036793 Homo Sapi
AZ602556 iM0421E20
A1267816 aq35c09 x
AU102743 AU102743
BEF013308 rk29c02.y
AA737998 nx15b01.s
AA737998 nx15b01.s
AA737998 nx15b01.s
AA737998 nx15b01.s
AZ774766 2M0004G05
AL048974 DKF2p566 x
AZ774766 2M0004G05
AL048974 DKF2p566 x
AL748117 Danio rer
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AL74817 Da
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AA853120 NHTBCae03
AJ499559 AJ499559
AU104458 AU104458
AU104458 AU104458
AU104922 AU104922
AA564185 nj0ddll.s
BI496942 df129910.
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AZ593107 1M0404022
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REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTTGGGTTTTAATTTTGTTTTATTGCACCTGACAAAATACAGTTATC 2932
                              Plate: 0206 row: D column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 47).

Dunn, D., Abyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ425831 47 bp DNA linear GSS 0: 1M0206D14F Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGCIM0206D14 F, DNA sequence.
                                                                                                                                                                                                                                                                                                     and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ425831
AZ425831.1
                                                                                    Pel: 801 585 5606
Pax: 801 585 7177

Camail: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                      University of Utah
University of Utah
                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                            Unpublished (2000)
                                                                                                                                                                                                                                                                                 plasmid inserts
                                                                                                                                                                                                                                                                                                 Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   house mouse
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Seq primer: Promega ~21m13
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This clone is available royalty
                                                                                                                                                                                     308,
quality sequence stop: 47
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              \(\text{riarmacia}\), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT/T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 20. Library constructed by Bento Soares and M.Fatima Bonaldo."
                                                                                                                                                                                 Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'note="Organ: breast; Vector: pT7T3D (Pharmacia) with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'clone_lib="Soares breast 3NbHBst"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:9606"
/clone="IMAGE:182126"
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                                                                                                                                                                                                                          Genome
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                                                                                                                                                                                                                                                                                             scaffolding with paired
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Pred. No. 7
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Best Local Similarity
                                                                                                                                                                                                 Clone from S. Wiemann, sequenced by sequencing consortium of the German
                                                           Berlin-Charlottenburg,
                                                                                             This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, H
                                                                                                                                                                                                                                                                                                          Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                       No s1 sequence available
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AL037866.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2784 GGCCAGCCTGTTTCTCTAACAAACAAACAAACAAACAACAGC
                                                                                                                                                                                                                                                                                                                                                                                                                  Bloecker H., Boecher M., Brandt P., Mewes W., Gassenhuber J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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12-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens mRNA;
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1 2 9 24 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were lighted to the blunt old in bith oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
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/db_xref="taxon:10090"
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Last updated, Version 1)
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Pred. No. 1.8e+05;
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                                                                                             Heubnerweg, 6, 14059
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSS
                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 40.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                  Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0152 row: I column: 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                   Plate: 0152 row: I column: 24
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="564 (synonym: hfbr2). Vectcr pAMP1; host
X1-2blue; sites NotI + SalI"
/dev_stage="fetal"
           (http://www.jax.org/resources/documents/dnares/). The DNP was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vertex. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/clone="DKFZp564J147"
                                                                                                                        Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC1M0152I24"
                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:9606"
adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                   /sex="Male"
                                                                                                                                                                                                                  /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                     Std Error:
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- COMMENT

TITLE

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2872 GTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATT 2909

Matches Query Match

28;

Conservative

0;

Mismatches

Indels

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Gaps

0;

Local

Similarity

0.7%; 73.7%;

Score 22; Pred. No.

No. 8.7e+05

DB 12;

Length 43;

ORIGIN

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VERSION
KEYWORDS
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         I (Dases 1 to -),
NIH-MGC http://mgc.nci.nih.gov/.
NIH-MGC http://mgc.nci.nih.gov/.
NIH-MGC http://mgc.nci.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
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                                                                                                                                                                                                                                 /Glone_IIAGE:333811"
/Glone_IIAGE:333811"
/Glone_IIAGE:74"
/Glone_IIAGE:74
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(where B = A, C, or G and N = A, C, G, or T). Average insert size 1.25 kb (range 0.6-4.0 kb). 14/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /db_xref-"taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism≖"Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%;
73.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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8.6e+05;
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RESULT 7
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KEYWORDS
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BJ047667/c
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       TITLE
                                                                                           AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                     2871 GGTTTTCTTGATGATTTTGGGTTTTAATTTTGTTTTTATTGCAC 2913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                     45
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                49 bp ruk62a04.y1 Schiller mouse MAC16 Mus IMAGE:1973550 5', mRNA sequence.
                        Marra,M., Hillier,L., Allen,M., Bowles,M., Dietric Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., M Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Theising,B., Wylle,T., Lennon,G., Soares,B., Wilso Waterston,R.
                                                                                                                                                                                                EST
                                                                                                                                                                                                                                                                                                                                                                                                                                        29;
       The WashU-HHMI Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BJ047667 NIBB Mochii normalized Xenopus neurula
BJ047667 NIBB Mochii normalized Xenopus neurula
                                                                                                                             Mammalia;
                                                                                                                                         Eukaryota; Metazoa;
                                                                                                                                                            Mus musculus
                                                                                                                                                                              house mouse.
                                                                                                                                                                                                                AI316598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
Tel: 81-559-81-6856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Tadasu Shin-i
Center For Genetic Resource Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Expressed genes in X. laevis embryo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                African clawed frog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kitayama,A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xenopus laevis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Xenopodinae; Xenopus.
                                                                                                      (bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        35
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: pBSRN3; Site_1: NotI; Site_2: EcoRI; cDNAs were oligo-dT primed and directionally cloned. Staging according to Nieuwkoop and Faber. Library is subtracted and was constructed by N. Garrett and A.M. Zorn, (Wellcome/CRC Institute)."
                                                                                                                         Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Wellcome/CRC Institute). "
4 c 3 g 3 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:8355"
/clone="XL017e20"
                                                                                                           1 to 49)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dev_stage="stage 15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'tissue_type="whole embryo"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Xenopus laevis"
                                                                                                                                                                                                                GI:4031865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Terasakà, C., Mochii, M., Ueno, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lib="NIBB Mochii normalized
                                                                                                                         Chordata;
Rodentia;
     EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 21.8; DB 13;
Pred. No. 1e+06;
0; Mismatches 14;
                                                                                                                      Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                      Bowles, M., Dietrich, N., Dubuque, T.,
                                                                                                                                                                                                                                                                            mRNA
                                                                                                                                                                                                                                                             musculus
                                                                      Martin, J., Morris, M.,
                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    others
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                                        Wilson, R: and
                                                                                                                                                                                                                                                             cDNA clone
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                                                                                                                                                                                                                                                                            EST 17-DEC-1998
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; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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AUTHORS
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AZ593107/c
LOCUS
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ORGANISM
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                                                                                                                                                                                                                        TITLE
                                                                                                                                                                                          JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2872 GTTTTCTTGATGATTTGGGTTTTAATTTTGTTTTTATTGCACCTGACAA 2920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0404 row: O column: 22
                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                      University of Utah
University of Utah
                                                                                                                                                                       Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                      Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                      1 (bases 1 to 49)
Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                              1M0404022F Mouse 10kb plasmid UUGC1M library clone UUGC1M04040402 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     32; Conservative
                                                                                                      84112, USA
                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end
                                                                                                                                                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                AZ593107.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ593107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicinep Washington University School of Medicinep Washington Darkway, Box 8501, St. L.
Seq primer:
                                                                                                                                                                                                                                                                                                                             Mammalia;
                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: -40RP from Gibco
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MGI:990290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact:
                                                                                                                      308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        40
                                                                                                                    Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University). "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               was constructed by Dr. Martin Schiller (Johns Hopkins
                                                                                                                                                                                                                                                                                                                             Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:1973550"
/clone_lib="Schiller mouse MAC16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/db_xref="taxon;10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /cell_line="colon cancer cell line"
/lab_host="SOLR"
 CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                GI:11715297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%;
                                                                                                                                                       Genome Center
                                                                                                                                                                       Weiss
                                                                                                                                                                                                                                                                                                                           Chordata;
Rodentia;
                                                                                                                    Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                           Barber, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 21.8; DB Pred. No. le+06; Mismatches
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Duval,B.,

Hamil,

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musculus genomic

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Length 49 Indels

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RESULT 9
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ORIGIN
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Best Local Similarity 65...
Matches 32; Conservative
                                      Clone from S. Wiemann, sequenced by sequencing consortium of the German No s1 sequence available
                                                                                                                  Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2803 CAAACAAACAAACAACAGCCTTGTTTCTCTAGTCACATCATGTGTATA 2851
  This clone is available at the RZPD in Berl: Please contact the RZPD: Ressourcenzentrum,
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens mRNA;
                                                                                                                                                                                                                                                                                                                                                                                                              12-MAR-1999
12-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AL039140.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AL039140;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSM003616
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                                                                                                                                                                          Bloecker H., Boecher M., Brandt P.,
                                                                                                                                                                                                                                                                                           Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                               EST; expressed sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49 CACACACACACACAGAGTTTTGCATGAAGTTATGTAATGTGTACA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
High quality sequent
                                                                                                                Klopferspitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                          (Rel. 59, Created)
(Rel. 59, Last updated,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (g1|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
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/clone="UUGC1M0404022"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                       EST DKFZp566L194_r1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence stop: 49
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Pred. No. 1e+06;
0; Mismatches ]
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                                                                                                                                                                                                                                                                                                                                                                                                              Version 1)
                                                                            GBF within the
                                                                                                                                                                      Mewes W., Gassenhuber J.,
                                                          Genome Project
                        Berlin
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Heubnerweg 6, 14059
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                                                                                                                                                                                                                                                              source
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                                                                                                                                                                                                                                                                                                                                                 Contact: Hideko Urushihara
Institute of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                Takeuchi,I., Kohara,Y. and Tanaka,Y.
Population analysis of CDNAs from unicellular and multicellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dictyostelium discoideum
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                                                                                                                                                                                                                                                                                                                                                                                        stages or
                                                                                                                                                                                                                                                                                                                                                                                                                          Urushihara,H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dictyostelium discoideum.
                                                                                                            Similarity
                                                                                                                                                                                                                                                                                     Email: hideko@biol.tsukuba.ac.jp
                                                                                                                                                                                                                                                                                                                        1-1-1 Tennoudai, Tsukuba, Ibaraki 305-8572,
                                                                                                                                                                                                                                                                                                                                      University of Tsukuba
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xl-2blue; sites NotI + SalI"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/clone="DKFZp566L194"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                          /dev_stage="vegetative"
3 c 1 g
                                                                                                                                                                                                             /clone="VSJ664"
                                                                                                                                                                                                                         /strain="AX4"
/db_xref="taxon:44689"
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                                                                                                                                                                                                /clone_lib="VS"
                                                                                                                                                                                                                                               organism="Dictyostelium discoideum"
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                                                                                                             Score 21.4; DB 9;
Pred. No. 1.3e+06;
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DEFINITION Locus

AZ310534 43 bp DNA linear GSS 29-SEP-20 1M0025J03R Mouse 10kb plasmid UUGCIM library Mus musculus genomic

29-SEP-2000

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RESULT 12
AL587842
LOCUS
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                                                                                                                                                         2801 AACAAACAAACAAACAA
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   AL587842
AL587842 BP Chicken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0025 row: J column: 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb
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AZ310534
AZ310534.1 GI:10352617
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                              10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|afi129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                            adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
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/clone="UUGC1M0025J03"
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/strain="C57BL/6J"
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100.0%;
44 bp mRNA linea
Brain Library Gallus gallus
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1.6e+06;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                            36 bp DNA linear GSS 05 1M0358G17F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0358G17 F, DNA sequence.
                  University of Utah Genome Center University of Utah
                                                                                                                         and Wright, D., Weiss, R.
                                                                                                                                                                                                                  Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 36)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27;
                                                           Contact: Robert B.
                                                                            Unpublished (2000)
                                                                                                   plasmid inserts
                                                                                                                   Mouse whole genome
                                                                                                                                                                                                  Dunn, D., Aoyagi, A., Barber, M.,
                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                          Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
                                                                                                                                                                                                                                                                                                     house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: frazer.murray@bbsrc.ac.uk
GCGGCCGCTTTTTTTTTTTTTTTTT 3' Poly A RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Roslin, Midlothian, EH25 9PS, Tel: +44 (0)131 527 4200 Fax: +44 (0)131 440 0434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chicken.
Gallus gallus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP Chicken Brain Library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Murray, F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ROS064C05, mRNA sequence AL587842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: M13F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Vector: pSPORT1; Site_1: Not1; Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT. 5' adaptor sequence: 5' TGACCTCGAG 3', 3' adaptor sequence: 5' GCGCCCCTTTTTTTTTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clonetech (*6854-1)"
0 c 4 g
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/dev_stage="Unknown"
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Pred. No. 1.6e+06;
0; Mismatches 10
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Sciurognathi; Muridae; Murinae; Mus
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Email:

ddunn@genetics.utah.edu

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Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens mRNA; EST DKFZp564K0763_r1
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                                                                                                   Duesterhoeft A., Lauber J., Mewes W., Gassenhuber J.,
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12-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expressed sequence tag
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Seq primer: CGTTGTAAAACGACGGCCAGT
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/strain="C57BL/6J"
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lab_host="E. Coli strain XL10-Gold, Ti-resistant, F
חייי בייי בייי חייים און Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 selected for ampicillin resistance.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACCESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                            JOURNAL
                                                                                                                                                                                                                                                                                                                                         TITLE
                                                                                                                                                                                                                                                                                                                                                                                                 AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This clone is available at the RZPD in Berlin Please contact the RZPD: Ressourcenzentrum, Heubnerweg Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 44 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequencing consortium of the German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clone from S. Wiemann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MIPS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     No si sequence available
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Am Klopferspitz 18a D-82152 Martinsried, GERMANY
                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Bar
Islam,H., Longacre,S.,
,M., Rose,M., Rose,R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ602556
45 bp DNA linear GS: 13-DEC-200 1M0421E20F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0421E20 F, DNA sequence.
                                                                                                 High quality sequence stop: 45
Location/Qualifiers
                                                                                                                                           Plate: 0421 row: E column: 20 Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                University of Utah
University of Utah
                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                        plasmid inserts
                                                                                                                                                                                                                                                                                                                                      Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                      and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ602556
                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                lass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 45)
                                                                                                                                                                                                                                                    308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 A; 6 C; 10 G; 4 T; 3 other;
                           /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0421E20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              x1-2blue; sites NotI + SalI"
/dev_stage="fet=""
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/clone="DKFZp564K0763"
/clone_lib="564 (synonym:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                     organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GI:11724746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequenced by
                                                                                                                                                                                                                                                                              Genome
                                                                                                                                                                                                                                                                                             Weiss
                                                                                                                                                                                                                                                                                                                                      scaffolding with paired
                                                                                                                                                                                                                                                                                                                                                                                               Barber, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                         Std Error:
                                                                                                                                                                                                                                                                                                                                                                   Stokes, R.,
                                                                                                                                                                                                                                                                                                                                                                   Mahmoud,M., Meenen,E., Pedersen,T., Reilly
Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                               Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No. 1.8e+06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Qiagen within the cDNA Genome Project
                                                                                                                                                                                                                                                                                                                                                                                               Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                           0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                    end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vector pampl; host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                  20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6
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Coli strain XL10-Gold, T1-resistant,

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BASE COUNT
ORIGIN
Search completed: May 16, 2003, 10:26:05
Job time : 4055 secs
                                                                                                                                                                                                                                    Query Match 0.7
Best Local Similarity 91.7
Matches 22; Conservative
                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                0.7%;
                                                                                                                                                                                                                                    Score 20.8; DB 17; Pred. No. 1.9e+06; 0; Mismatches 2;
                                                                                                                                                                                                                                                                                             Length 45;
                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                       0,
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